

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: May 11, 2004, 15:24:27 ; Search time 18.8426 Seconds

(without alignments)
2295.997 Million cell updates/sec

Title: US-09-927-315-9

Perfect score: 4443
Sequence: 1 MGPRAKTICSLFLLMWLAE.....ERNTPAYNSMIGYTRRD 838

Scoring table:

BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Database :

Issued Patents AA:*
1: /cgn2_6/ptodata/2/1aa/5A_COMB.pep:*
2: /cgn2_6/ptodata/2/1aa/5B_COMB.pep:*
3: /cgn2_6/ptodata/2/1aa/6A_COMB.pep:*
4: /cgn2_6/ptodata/2/1aa/6B_COMB.pep:*
5: /cgn2_6/ptodata/2/1aa/6C_COMB.pep:*
6: /cgn2_6/ptodata/2/1aa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	3514	79.1	669	4	US-09-361-631-7
2	3203.5	72.1	843	4	US-09-361-631-1
3	3151.5	70.9	843	4	US-09-361-631-2
4	1107	24.9	1059	3	US-09-134-513-2
5	1094	24.6	1078	1	US-08-485-588-7
6	1094	24.6	1078	1	US-08-484-565-7
7	1094	24.6	1078	2	US-08-480-751-7
8	1094	24.6	1078	2	US-08-943-986-7
9	1094	24.6	1078	3	US-08-353-784-7
10	1094	24.6	1078	3	US-08-484-719B-7
11	1094	24.6	1078	4	US-08-484-159-7
12	1092.5	24.6	1027	4	US-09-162-021B-2
13	1088.5	24.5	1079	4	US-08-485-588-8
14	1088.5	24.5	1079	1	US-08-484-565-8
15	1088.5	24.5	1079	2	US-08-480-751-8
16	1088.5	24.5	1079	2	US-08-943-986-8
17	1088.5	24.5	1079	3	US-08-353-784-8
18	1088.5	24.5	1079	3	US-08-484-719B-8
19	1088.5	24.5	1079	4	US-08-484-159-8
20	1087.5	24.5	1085	1	US-08-485-588-5
21	1087.5	24.5	1085	1	US-08-484-565-5
22	1087.5	24.5	1085	2	US-08-480-751-5
23	1087.5	24.5	1085	2	US-08-943-986-5
24	1087.5	24.5	1085	3	US-08-353-784-5
25	1087.5	24.5	1085	3	US-08-484-719B-5
26	1087.5	24.5	1085	4	US-08-484-159-5
27	1079	24.3	1088	1	US-08-485-588-6

28	1079	24.3	1088	1	US-08-484-565-6	Sequence 6, Appl1
29	1079	24.3	1088	2	US-08-480-751-6	Sequence 6, Appl1
30	1079	24.3	1088	2	US-08-943-986-6	Sequence 6, Appl1
31	1079	24.3	1088	3	US-08-353-784-6	Sequence 6, Appl1
32	1079	24.3	1088	3	US-08-484-719B-6	Sequence 6, Appl1
33	1079	24.3	1088	4	US-08-484-159-6	Sequence 6, Appl1
34	1012.5	22.8	877	4	US-09-619-353-2	Sequence 2, Appl1
35	995	22.4	1219	2	US-08-687-289A-6	Sequence 6, Appl1
36	995	22.4	1219	4	US-09-435-887-6	Sequence 6, Appl1
37	964	21.7	975	4	US-09-695-481-4	Sequence 4, Appl1
38	917.5	20.7	863	4	US-09-619-353-14	Sequence 14, Appl1
39	867.5	19.5	851	4	US-09-619-353-12	Sequence 12, Appl1
40	856	19.3	854	4	US-09-619-353-10	Sequence 10, Appl1
41	850	19.1	856	4	US-09-619-353-8	Sequence 8, Appl1
42	849.5	19.1	835	4	US-09-619-353-7	Sequence 7, Appl1
43	743	16.7	1058	2	US-08-687-289A-5	Sequence 5, Appl1
44	743	16.7	1058	4	US-09-435-887-5	Sequence 5, Appl1
45	724.5	16.3	915	1	US-08-453-862-2	Sequence 2, Appl1

ALIGNMENTS

```
RESULT 1
US-09-361-631-7
; Sequence 7, Application US/09361631
; Patent No. 6383778
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliot
; APPLICANT: Lindemeyer, Uergeren
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; FILE REFERENCE: 02307B-088720US
; CURRENT APPLICATION NUMBER: US/09/361,631
; EARLIER FILING DATE: 1999-07-27
; EARLIER APPLICATION NUMBER: US 60/095,464
; EARLIER FILING DATE: 1998-07-28
; EARLIER APPLICATION NUMBER: US 60/112,747
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 7
; LENGTH: 669
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: human G-protein coupled receptor (GPCR) B4 amino
; OTHER INFORMATION: acid sequence
US-09-361-631-7
Query Match          79.1%; Score 3514; DB 4; Length 669;
Best Local Similarity 98.4%; Pred. No. 0;
Matches 666; Conservative 1; Mismatches 2; Indels 8; Gaps 2;

QY 162 ITTSAISSELDKRPALRTTSPADHVEAMQVMHFRNNWITIVSSPTGRDNGQ 221
DB 1 ITTSAISSELDKRPALRTTSPADHVEAMQVMHFRNNWITIVSSPTGRDNGQ 60
QY 222 LLGSEVARDICIAFOETPLTPQNMNTSESRORLVTVKLOOSTARVVVPSDILT 281
DB 61 LLGSEVARDICIAFOETPLTPQNMNTSESRORLVTVKLOOSTARVVVPSDILT 120
QY 282 YHFNVEVLKQFTGAVWIASSEMAIDPVNLHTELGHLGTFLGTTIOGVP1PGFSEPREW 341
DB 121 YHFNVEVLKQFTGAVWIASSEMAIDPVNLHTELGHLGTFLGTTIOGVP1PGFSEPREW 180
QY 342 GPQAGPPPLSTSGSYTNOECNCLNATLSFNTTLRSGERVVYSVSAVVAHALHS 401
DB 181 GPQAGPPPLSTSGSYTNOECNCLNATLSFNTTLRSGERVVYSVSAVVAHALHS 240
QY 402 LLGCKSKCTKRVVYPMQLLBEIKWNTLLDHOIFPDQGDVALHLLEIVQMWDKSGNP 461
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Db 241 LLGCKSTCTKRVVPMQLEBIEIKVNFLLDHOJFPDQGDVALHLEIVQMQRSONP 300
Qy 462 FOSVASYPLRQRLKIDISHTNTNTIPMSCKRQSQGQKKRPVCIHNCCEPIDCL 521
Db 301 FOSVASYPLRQRLKNIK-TSLHTVNTNTPMSCKRQSQGQKKRPVCIHNCCEPIDCL 359
Qy 522 PGTPLNTHEDVEYECQACPNNEWSYQSESCFQRQLVFLMEHAPITAAVLAALGFLSTL 581
Db 360 PGTPLNTHEDVEYECQACPNNEWSYQSESCFQRQLVFLMEHAPITAAVLAALGFLSTL 412
Qy 582 AILVIFWHRFQTPYRSAGGPMCFMLTLVAVVWVYVYVGPVAVSTCLCRQALPLCF 641
Db 413 AILVIFWHRFQTPYRSAGGPMCFMLTLVAVVWVYVYVGPVAVSTCLCRQALPLCF 472
Qy 642 TICISCIARVSFOIVCAFKMSRPRAYSVMYRQGVPMNAFTVLKXVIVIGMLARP 701
Db 473 TICISCIARVSFOIVCAFKMSRPRAYSVMYRQGVPMNAFTVLKXVIVIGMLARP 532
Qy 702 QSHRTDDBDKITIVSCNPNYRNSLTFNTSLDLLSVGFSFAYMGKELPTNYNEAKFI 761
Db 533 QSHRTDDBDKITIVSCNPNYRNSLTFNTSLDLLSVGFSFAYMGKELPTNYNEAKFI 592
Qy 762 TLSMTFYFTSSVSLCTFMSAYSGVLVTIVDLLVTLNLALISLGYFGPKCYMILFYPERN 821
Db 593 TLSMTFYFTSSVSLCTFMSAYSGVLVTIVDLLVTLNLALISLGYFGPKCYMILFYPERN 652
Qy 822 TPAYNSMIOGYTMRD 838
Db 653 TPAYNSMIOGYTMRD 669

RESULT 2

US-09-361-631-1
; Sequence 1, Application US/09361631
; Patent No. 6383778
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliot
; APPLICANT: Lindemeier, Juergen
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; FILE REFERENCE: 02307E-088720US
; CURRENT APPLICATION NUMBER: US/09/361,631
; EARLIER FILING DATE: 1999-07-27
; EARLIER APPLICATION NUMBER: US 60/095,464
; EARLIER FILING DATE: 1998-07-28
; EARLIER APPLICATION NUMBER: US 60/112,747
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURES:
; OTHER INFORMATION: rat G-protein coupled receptor (GPCR) B4 amino
; OTHER INFORMATION: acid sequence
US-09-361-631-1

Query Match 72.1%; Score 3203.5; DB 4; Length 843;
Best Local Similarity 70.6%; Pred. No. 6.2e-296;
Matches 595; Conservative 108; Mismatches 133; Indels 7; Gaps 4;
Qy 1 MGPAKTIQSLFLLIWLAE--AENSDFYLPQGYLLGLPSLHANKGIVHLNFIQVP 57
Db 1 MGPAKTIQSLFLLIWLAE--AENSDFYLPQGYLLGLPSLHANKGIVHLNFIQVP 60
Qy 58 MCKEYEVVIGVYVNMQAMRFAVEEINNDSSLLPGULGYELVDVCYISNNQPVLYPLAH 117
Db 61 KCNFEYTKVGLVYVNMQAMRFAVEEINNDSSLLPGVLLGYEMVDVCYISNNIHPIGLYPLAQ 120

Qy 118 EDNLPIQEDYSNYSIRVAVVIGPDNSESVMYANFLSLPQLQITYSASIDELRDKVR 177
Db 121 DDLLPILKQYQVMRPHVAVIGPDNSESALITYSNLSHFLPQITYSASIDELRDKVR 180
Qy 178 PALRTTPSADHVEANVQMLHFRKMWIIVLVSSDPTTYGNDGOLLGERAR-RDICIAP 236
Db 181 PSMRLTPSADHVEANVQMLHFRKMWIIVLVSSDPTTYGNDGOLLGERAR-RDICIAP 240
Qy 237 QETPLTQPNQNMSESRORLVTVLIDLOOSTARVVVVFSPDLTLVHFNEVLQRNFTGA 296
Db 241 QETPLTQPNQNMSESRORLVTVLIDLOOSTARVVVVFSPDLTLVHFNEVLQRNFTGA 300
Qy 297 VMASEMAIDPVLANLTTELGLTFLGIIQSVPIRPFSEFRMGQAGPPLSRTSQS 356
Db 301 VMASEMAIDPVLANLTTELGLTFLGIIQSVPIRPFSEFRMGQAGPPLSRTSQS 360
Qy 357 YTCNQEDNCINATLSNTILRLSGERVVYSVSAVAVAAHLSLGCDSKSTTKRVY 416
Db 361 YTCNQEDNCINATLSNTILRLSGERVVYSVSAVAVAAHLSLGCDSKSTTKRVY 420
Qy 417 PMOLLEIKVNFLLDHOJFPDQGDVALHLEIVQMQRSONPQOSVASYPLRQRLK 476
Db 421 PMOLLEIKVNFLLDHOJFPDQGDVALHLEIVQMQRSONPQOSVASYPLRQRLK 480
Qy 477 NIODISHTVNTNTPMSCKRQSQGQKKRPVCIHNCCEPIDCLPGTPLNTHEDVEYECQ 536
Db 481 YINNVSMYTPNNTVPVMSCKRQSQGQKKRPVCIHNCCEPIDCLPGTPLNTHEDVEYECQ 540
Qy 537 ACNNNEWSYQSESCFQRQLVFLMEHAPITAAVLAALGFLSTLAILVIFWHRFQTPY 596
Db 541 ACNNNEWSYQSESCFQRQLVFLMEHAPITAAVLAALGFLSTLAILVIFWHRFQTPY 600
Qy 597 RSAGGPMCFMLTLVAVVWVYVYVGPVAVSTCLCRQALPLCFITCISCIARVSFOIV 656
Db 601 RSAGGPMCFMLTLVAVVWVYVYVGPVAVSTCLCRQALPLCFITCISCIARVSFOIV 660
Qy 657 CAFQMSRPRAYSVMYRQGVPMNAFTVLKXVIVIGMLARPQSH--RTDPPDDPKI 714
Db 661 CAFQMSRPRAYSVMYRQGVPMNAFTVLKXVIVIGMLARPQSH--RTDPPDDPKI 719
Qy 715 TIVSCNPNYRNSLTFNTSLDLLSVGFSFAYMGKELPTNYNEAKFTLSMTFYFTSSVS 774
Db 720 TIVSCNPNYRNSLTFNTSLDLLSVGFSFAYMGKELPTNYNEAKFTLSMTFYFTSSVS 779
Qy 775 LCTFMSAYSGVLVTIVDLLVTLNLALISLGYFGPKCYMILFYPERNTPAYNSMIOGYT 834
Db 780 LCTFMSAYSGVLVTIVDLLVTLNLALISLGYFGPKCYMILFYPERNTPAYNSMIOGYT 839
Qy 835 MRR 837
Db 840 MRR 842

RESULT 3

US-09-361-631-2
; Sequence 2, Application US/09361631
; Patent No. 6383778
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliot
; APPLICANT: Lindemeier, Juergen
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; FILE REFERENCE: 02307E-088720US
; CURRENT APPLICATION NUMBER: US/09/361,631
; EARLIER FILING DATE: 1999-07-27
; EARLIER APPLICATION NUMBER: US 60/095,464
; EARLIER FILING DATE: 1998-07-28
; EARLIER APPLICATION NUMBER: US 60/112,747
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0

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SEQ ID NO 2
LENGTH: 843
TYPE: prt
ORGANISM: Mus sp.
FEATURE:
OTHER INFORMATION: mouse G-protein coupled receptor (GPCR) B4 amino
OTHER INFORMATION: acid sequence
US-09-361-631-2

Query Match      70.9%; Score 3151.5; DB 4; Length 843;
Best Local Similarity 68.9%; Pred. No. 5.5e-291;
Matches 581; Conservative 113; Mismatches 142; Indels 7; Gaps 4;

QY 1 MGPRAKTIQSLFLLVLAEP---AENSDFLPDQVLLGSLSLHANMKGIYHLNFIQVP 57
DB 1 MGPQATTLHLPLLLHALPKPVMVLVGNSDPLAGDGLPTLLANVKSLSLTLQVP 60
QY 58 MCKEYEVKIGYNLMQAMRFAVEEINNDSSLPGVLLGEYIVDVCYSNNQPVLYFLAH 117
DB 61 KCEYEMKVLGYNLQAMRFAVEEINNCSSLLPGVLLGEMVDVCLSNIQPGLYFLSQ 120
QY 118 EDNLLPIQEDYSNYSRVVAIVIGPDNSSEWTVANFLSLPQITYSASIDELADKVF 177
DB 121 IDDFLPILKDYQYRQVAVAVIGPDNSSESAITVSNILSYFLVPQVYSAITDKLDKRF 180
QY 178 PALLETTSADHHEAVMQLMHPFRNMIIVLVSSDTYGRDNGQLLGERVARR-DICIAF 236
DB 181 PMLKTPVPAATHIEAMQVMVHFMVNIIVLVSDDYGRNSHLSQRLTNTGDIQIAF 240
QY 237 QETLPTLPQNMWTSERQRLVTIVDKLOOSTARVVVVSPDLTYHFNENEVLROQFTGA 296
DB 241 QEVLVPERNOAVRREBOQDNIDKLRISARVVIVISPELSLHNFREVLKRNFTCF 300
QY 297 WIASESMAIDVNLHNLTELGLTFLGTTIGSVPIPGSESEFREMGPQAPPLSRTSQS 356
DB 301 WIASESMAIDVNLHNLTELRLHTGTFLGTVIGRVSIPGSGRVRHDKRGMPMETSIR 360
QY 357 YTCNQCNCNLAATLSFNTILSGRNVYYSVAVYANAHLSLGDCKSTCTKRVY 416
DB 361 TTCNODCDACMNTISFNNVLMHSGRRVYVSAYAAVHLLHRLHCNQVRCQIY 420
QY 417 PWQLLEIWKVFTLLDHOIFPDQGVNLALEIVOMQDRSQNPQSVASVYPIOROLK 476
DB 421 PWQLLEIWKVFTLLDHOIFPDQGVNLALEIVOMQDRSQNPQSVASVYPIOROLK 480
QY 477 NIQDISMHTVNTIIPMSCKSKQSGQKKKPVGIIHVCEECIDCLPGTFLNHTEDYECQ 536
DB 481 YISNVSMTYNNVTIIPMSCKSKQSGQKKKPIGLHPCFECVDCPDDTYLNSVDENCL 540
QY 537 ACPNNEMSYQSTSCFKQQLVLEHHEATIAVALLAAGELSTALIVIFRHHQTPYV 596
DB 541 SCPGSMWSYKNNIACFKRLAFLHWEHEVETIIVTLLAAGFISTALILIFRHHQTPV 600
QY 597 RSAGSPMGLMTLLLVAVMVVYVGPVKVSTCROALPFLCFITCISCTAVSPQV 656
DB 601 RSAGSPMGLMTLLLVAVMVVYVGPVKVSTCROALPFLCFITCISCTAVSPQV 660
QY 657 CAFKMASRPPRAYSYWVRVYQGPVSMATITVLMVIVYIGMLARQDSHP--RTDDEPKI 714
DB 661 CVFKMARRLPSAYGEMRHYGPFVAFITAVKVALVAGNMMLA--TTINFIGRTPDDPMI 719
QY 715 TIVSCNPNYRNLRTSISDLLSVYGSFAYMGKELPNNYNEAKFITLSMFTYTSYS 774
DB 720 ILSCHPNYRNLRTSISDLLSVYGSFAYMGKELPNNYNEAKFITLSMFTYTSYS 779
QY 775 LCTFMSAVSGVLVTVDLLVTLNLAISLGFGPKCYMILFYPERNTPAYNSNIQGT 834
DB 780 LCTFMSAVSGVLVTVDLLVTLNLAISLGFGPKCYMILFYPERNTPAYNSNIQGT 839
QY 835 MKR 837
DB 840 MKR 842
```

```
RESULT 4
US-09-134-513-2
Sequence 2, Application US/09134513
Patent No. 6210964
GENERAL INFORMATION:
APPLICANT: Brown, Edward M.
APPLICANT: Diaz, Ruben
APPLICANT: Bai, Mei
TITLE OF INVENTION: The Avian Extracellular Calcium-Sensing
RECEPOTR
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Vinson & Elkins L.L.P.
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/134, 513
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/13003
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6585
TELEFAX: (202) 639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1059 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-09-134-513-2

Query Match      24.9%; Score 1107; DB 3; Length 1059;
Best Local Similarity 31.9%; Pred. No. 4.7e-96;
Matches 284; Conservative 148; Mismatches 354; Indels 104; Gaps 24;

QY 9 CSLFPL--WVLAEPENSDFYLPQDYLLGSLFSLHANMKGIYHLNFIQVPMCKEYEVK 66
DB 6 CCLILLLFTWNTAAAGPQQAQKKDITLGLFPIHF--GVAARD--QDLKSRDESEVC 60
QY 67 IGYNL-----NQAMFAVEEINNDSSLPGVLLGEYIYDVC-YISNNQPVLYFLA-HED 119
DB 61 IRYNFRGRMLQAMFAVEEINNSPMLPNMTLGRIPDTCTVSKALEATLSFVAQNKI 120
QY 120 NLLPIQ--DYSNYSRVVAIVIGPDNSSEWTVANFLSLPQITYSASIDELADKVF 177
DB 121 DSNLNDERCENSEHPTIATVAVGAGSVSTAVANLLGLFIIPQVSYASSRSLSNQOF 180
QY 178 PALLETTSADHHEAVMQLMHPFRNMIIVLVSSDTYGRDNGQLLGERVARRDICIAFQ 237
DB 181 KSFLETTINDEHQATAMADIIEYFRMNVGTIADDDYGRGIEKFRSEAESEEDICIDS 240
QY 238 ETLPTLPQNMWTSERQRLVTIVDKLOOSTARVVVVSPDLTYHFNENEVLROQFTGA 297
DB 241 ELI-----SQYSDEEILQVVVEI--QNSTARVIVVSSGPDLEPLIKELVRRNITKI 292
QY 298 WIASESMAIDVNLHNLTELGLTFLGTTIGSVPIPGSESEFREMGPQAPPLSRTSQS 357
DB 293 WIASESMASSSLIAMPEFRVIGTIGPALAAQOIGRFRFLQ--KVHPKKSANNGPAK 349
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358 TCNOBCDNC-----NATLSFN----- 374
350 EFWETFCIYLPSESKSPASAFKABEGGAGNGTAAPRPTGENTISVETPYMD 409
375 -TIRLSGERVYVSYAVAVAAHLSLGC-----DKSTCKRVYVPMOLLEBIW 425
410 FTHLRIS-----YNYVLAIVSIAHALODIYTCTPGKLFTHGSCADIKKVEAMQVKKILR 464
426 KVNFTL-IDHOFPPQGDVALHLEIVOMQDRSQNP--FQSVASYPL--OROLKNI 478
465 HLNFTSNNGEVDFFEDLVGNYSIIWMHLSPEDGSVFFEBGVHNYAKKGERLFTNE 524
479 ODISHVNTNTIPMSCKRCSGOKKKPV-GIHVCEFCIDCLPGTFLNTHTEDEYEQCA 537
525 NKILMSGSKVFPENCSDRCLPGRKGIIEGEPCCFECVDCPGEVSDET-DASADK 583
538 CPNNEMSYOSETSCPKQVLPLEMHEAPLIAVALLAALGFLSTALIVIPMRHFTPIVR 597
584 CPEDWYSENHTSCIPKQIEFLSMTEPFGIALTLFAVLGIFLTSFVLGVTKFRTPYIK 643
598 SAGEMCLMTLLLVAMVVPVYVPPKVSCTCROALLPLCTITICISIAVSFOIVC 657
644 ATNELSLYLLFSLCCFSSSLFFIGEPQNMTCRLRQAFGISFVLCISCLVKTNRVLL 703
658 AFKMSRPPRA-VSYVWRYQPVYMAFITYLVKVIYVIGMLARPOSHPRTPDDPKITI 716
704 VFE-AKPTSLHRKMGMLNQLFVLVPLCTPQIYICIMLYTAPSSYRNHELEDEIIF 761
717 VSCNPNVNSLLFNTSLDLLSVGSPFAYMGKELPTVYNEAKFTTSMTEFTSSVSLC 776
762 ITCHEGSLMALGFLGYTCLLAICFPFAFKSRKLPENFNEXAKFTTFMILFFIWMIS-- 819
777 TFMASGVLTIVDLYTVNLALISIG----YFGRPCVMTLFPRENT 822
820 -FIPAYASTYKQFVS-AVEVIAIILASFGLLACIFFNKVYIILFPRSHNT 867

RESULT 5

US-08-485-588-7
Sequence 7, Application US/08485588
Patent No. 5688938

GENERAL INFORMATION:
APPLICANT: Edward M. Brown

APPLICANT: Steven C. Hebert

APPLICANT: Forrest H. Puller

APPLICANT: James E. Garrett, Jr.

TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE

NUMBER OF SEQUENCES: 20

CORRESPONDENCE ADDRESS:

ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA

ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PASTSEO

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,588

FILING DATE: 7 June, 1995

CLASSIFICATION: 435

PRIOR APPLICATION DATA: including application

PRIOR APPLICATION DATA: described below: 9

APPLICATION NUMBER: 08/353,784

FILING DATE: 9 December, 1994

FILING DATE: 9 December, 1994

APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Hebert, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-485-588-7

Query Match 24.6%; Score 1094; DB 1; Length 1078;
Best Local Similarity 31.5%; Pred. No. 8.4e-95;
Matches 279; Conservative 160; Mismatches 342; Indels 106; Gaps 26;

16 WVLPAENSDPYLP-----GDYLLGGLFSLHANKGIHNLQVMPCKEYEVYIG 68
8 WVLLALTMHTSAYGPDQRAQKGDIIIGLFPPIHF--GVAAXD--QDLKSRESVACIR 62
69 YNL-----MQMRPAVEINNDSSLPGLGLEYIVVC-YISNNVPVLYFLA-HEDNL 121
63 YNFRGFRMLQMTFAIEINSSPALPLNLTGYRIPTCTVTSVLEATISFVAQNTIDS 122
122 LPIDQ--DYSNYSRVAVAVIGPDNSSEVMTVAFNLSFLPLQITYSAISDELARDKVFPA 179
123 LNLDFCNCSEHISTTAIVGATSGSVTAVALGLFYIPQVSYASSRLSKNPKS 182
180 LLRTTSPADHHEAVMVLHFRNMWIIIVSSDTYGRDNGQLGERVARRDICIATQET 239
183 FLRTIPDEHQATAMADIIIEYFRNMVGTIAADDYGRPGIEKREBAERDIDIPSEL 242
240 LPTIQPNQMTSEERQVLYTVKLGQSTARVYVESPDLTVHFFNEVLKONTGAVMI 299
243 I-----SQYSDDEEIQHVEVI---QNSTAKVIVVFSSGPDLEPLIKEIVARNITGKWL 294
300 ASESMAIDPVLMNLTGLHGTFLGITIQSVPIPGFSEF-----REMGPO 344
295 ASEMASSSLIAMQYHVHVGITIGFALKAGQIDGFEFLKAKHPRKSVNNGFAKEWEE 354
345 A-----GPPRLTSQSYTCNQECNCLNATLSFNTILRSGE----- 382
355 TFNCHLQGAAGPLFVDTPLRGH--EESGDRFSSSAFAPPL--CTDENISSVETPYID 410
383 ----RVYYSYSAVYAVAAHLSLGC-----DKSTCKRVYVPMOLLEBIWKNFT 430
411 YTHLRISYNYVLAIVSIAHALODIYTCLPGKLFTHGSCADIKKVEAMQVKKILRHLNFT 470
431 -LIDHOFPPQGDVALHLEIVOMQDRSQNP--FQSVASYPL--OROLKNIODIS 482
471 NMGEQVTFDECGDLVGNYSIIWMHLSPEDGSIVFKV-GYVNYVYAKKGERLFINNEKIL 529
483 WHTVNTNTPMSCKRCSGOKKKPV-GIHVCEFCIDCLPGTFLNTHTEDEYEQACPPNN 541


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QY      720  NPNYRSLLENTSLDLSIVSGSPAYMKEKLEPTNYNEAKFTLSMTFYSVSLCTEM 779
          |||:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|
Db      766  HEGSLMALGTLGYTCLLAICFPFRKSKLPENFENAKPTTFEMLIFELWIS---FI 822
          |||:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::
QY      780  SAYSGVLTVTVLLVTVNLLAISLG---YGPCKYMLTFYPERNT 822
          |||:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::
Db      823  PAASATYGVFS-AVEVIAILAAFCLLACIFENNYIILLFEPSENNT 868
          |||:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::|:::

```

```

1  US-08-480-751-7
2  Sequence 7, Application US/08480751
3  Patent No. 5858684
4  GENERAL INFORMATION:
5  APPLICANT: Edward F. Nemeth
6  APPLICANT: Edward M. Brown
7  APPLICANT: Steven C. Hebert
8  APPLICANT: Forrest H. Fuller
9  APPLICANT: James E. Garrett, Jr.
10 TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
11 TITLE OF INVENTION: MOLECULS
12 NUMBER OF SEQUENCES: 20
13 CORRESPONDENCE ADDRESS:
14 ADDRESSEE: Lyon & Lyon
15 STREET: First Interstate World Center
16 STREET: Suite 4700
17 STREET: 633 West Fifth Street
18 CITY: Los Angeles
19 STATE: California
20 COUNTRY: USA
21 ZIP: 90071
22 COMPUTER READABLE FORM:
23 MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
24 COMPUTER: IBM PC compatible
25 OPERATING SYSTEM: PC-DOS/MS-DOS
26 SOFTWARE: FASTSEQ
27 CURRENT APPLICATION DATA:
28 APPLICATION NUMBER: US/08/480,751
29 FILING DATE: 7 June, 1995
30 CLASSIFICATION: 435
31 PRIOR APPLICATION DATA:
32 PRIOR APPLICATION DATA: including application
33 PRIOR APPLICATION DATA: described below: 9
34 APPLICATION NUMBER: 08/355,784
35 FILING DATE: 9 December, 1994
36 APPLICATION NUMBER: PCT/US/94/12117
37 FILING DATE: 21 October, 1994
38 APPLICATION NUMBER: U.S. 08/292,827
39 FILING DATE: 23 August, 1994
40 APPLICATION NUMBER: U.S. 08/141,248
41 FILING DATE: 22 October, 1993
42 APPLICATION NUMBER: U.S. 08/009,389
43 FILING DATE: 23 February, 1993
44 APPLICATION NUMBER: U.S. 08/017,127
45 FILING DATE: 12 February, 1993
46 APPLICATION NUMBER: U.S. 07/934,161
47 FILING DATE: 21 August, 1992
48 APPLICATION NUMBER: U.S. 07/834,044
49 FILING DATE: 11 February, 1992
50 APPLICATION NUMBER: U.S. 07/749,451
51 FILING DATE: 23 August, 1991
52 ATTORNEY/AGENT INFORMATION:
53 NAME: Heber, Sheldon O.
54 REGISTRATION NUMBER: 38,179
55 REFERENCE/DOCKET NUMBER: 213/004
56 TELECOMMUNICATION INFORMATION:
57 TELEPHONE: (213) 489-1600
58 TELEFAX: (213) 955-0440
59 TELEX: 67-3510
60 INFORMATION FOR SEQ ID NO: 7:
61 SEQUENCE CHARACTERISTICS:
62 LENGTH: 1078 amino acids
63 TYPE: amino acid

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;      TOPOLOGY: linear
;      MOLECULE TYPE: protein
US-08-480-751-7

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Query Match	24.6%	Score 1034	DB 2	Length 1078
Best Local Similarity	31.5%	Pred. No. 8.4e-95		
Matches 279	Conservative 160	Mismatches 342	Indels 106	Gaps 26

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0Y      16 WTLAEARENSDPFLP-----GDVLLGGTSLSAHNMGIYHNLFLQVPMCKEYEVYIG 68
0Y      8 WVLALATWHTSAAGPORAOKKGDITLGGFLPFHF--GVAAKO--DOLKSPRESVEICIR 62
0Y      69 YNL-----MOAKFAVEIEINDDSSLLPGLLGYEIVDVC-YISNNVOPLYFLA-HEBNL 121
0Y      63 YNRGRFRWLQAMIFALIEEINSSPALLPNLTGCRFPDCCNVSXALENTLSFVAKNIDS 122
0Y      122 LP1QE--DYSNYSIRVAVVIGPDNSSEWVTYANFLSLFLLPQITYSAISDELKDKVFP 179
0Y      123 LNFDECNCSSEHPISTIAVVGATGSGVSTAVANLLGLFPYIQVSVYASSRLLSKNQPKS 182
0Y      180 LLAFTTSADHHEAMQMLHFRNMWIVLVSDPYGGDNQQLGEEVARRDICIACQET 239
0Y      183 FLRTIENDEQATAMADIIIEFRNMWVGITIAADDYGGPPIEKFEAREAEEDDICIIDSEL 242
0Y      240 LPFLPONMNTSEERQRLVTIVDKQOSTARVVVFPDILLYHFNFEVLRONFTGAVMI 299
0Y      243 I-----SQSDEBEIOWVEVI-----QNSTAKYIVFSSGPRLEBLIEIRNRTGKMW 294
0Y      300 ASSSMAIDPVLNHLTELGLGTFLLGITTQSVAPIPGFSSE-----REWKPO 344
0Y      295 ASBAMASSSLIAMPQYFHVHVGCTIGFALKACQIPGFRFELKVKHPRKSVHNGFAKEFWEE 354
0Y      345 A-----GPPRLSRTSQSYTCNOQCDNCNATLSFNTILASGE-----362
0Y      355 TENCHLOEGAKGAPLDVDFLRGH--EESGDPRSNTSIAFRPL--CTGDENISVETPYID 410
0Y      383 -----RVVYSVYSAVVYVAHALHSLIGC-----DKSTCKRVVYPMOLLEIMKNFT 430
0Y      411 YTHLRISTYNYVLAIVSIALADODITCLPBGKGLPTNGSCADIKKVEAMQYUAKHLRIHFT 470
0Y      431 -LLDHQIFPDPOGVALHLEIVQWQMDBSQNP--FQSVASYRPL-----OROLKNIDIS 482
0Y      471 NNMGEQVTFDECGDVGANYSIIMHLSPEDEGSIVFEKEY-GYNNYVAKKGERLFINBEKIL 529
0Y      483 WHIVNNTIMSMNSKRCQSGCKKKRV-GIHNCCEPCIDCLPRTLHNTDEYEQACAPNN 541
0Y      530 WSGFSEVPSPNSCRCLAGTRKGIIEBPPTCCCEVCBPGEHSDEL-DASAKNCRPD 588
0Y      542 EMWQSETSCEFQKROLFTEHMEAPTAVALAALGFLSTLAIYFMRHFOPTPIVSAGG 601
0Y      589 FMENENHTSCIAKEIEFLSMTEPFCIALTLFRAVIGIFLTAVLGVFIKPNPTPIVAKTNR 648
0Y      602 PMCFMLTLTLVAIVVVYVGVPRPVSTCLCQALFPLCFTTICISCIASVBSFOICAF-- 659
0Y      649 ELSYLLLFLLCLCFSSSLFFIEBPQDMTCRLRQPAFGISFVLCISCIILVXTNRVLVFEA 708
0Y      660 KMSRPRRAYSVWRQSGPYVSMARITLAKNVIIVIGLARPOSHPRTPDDRPITIVSC 719
0Y      709 KILTSYHRK--WMGLNQFLVPLCTMQMIVICIMLYTAPBSYNOLEDEDIITTC 765
0Y      720 NPYVRSLFLFNTSLDLILSVGVFSFAYVNGKELPTVNEAKFITLSMTFYTSSVSLCTFM 779
0Y      766 HESLSMALGFLIGYTCLLAICFFPAFASRKLPENFNEAKITTSMLIFLIWIS--FI 822
0Y      780 SAYSGVLVTIVDLVTVNLALISIG---YFGPKCMIILFPERNT 822
0Y      823 PAVASTYGRFVS-AVEVIAIILASGLACIFFNKIYIILFKPSRNT 868

```

GENERAL INFORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESS: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/943,986
FILING DATE: 03-OCT-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/484,565
FILING DATE: 7-June-1995
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/006
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-943-986-7

Query Match 24.6%; Score 1094; DB 2; Length 1078;
Best Local Similarity 31.5%; Pred. No. 8.4e-95;
Matches 279; Conservative 160; Mismatches 342; Indels 106; Gaps 26;

QY 16 WVLAEPAENSDPYLPE-----GDVLLGLFSLHANKGIYVLTNLOYVPMCKEYEVKYG 68
DB 8 WVLALLTHTSTAYGDDQAKKKDITLLGLPFIHF--GVAKD--QDLKRPSEVECTR 62

QY 69 YNL-----MQAMRFVVEEINDDSLLPVLLGLEYIVDVC-YISNNVQVLYFLA-HEENL 121
DB 63 YNFGFWLQMIIFAEIEINSSPALLPMLTLGYRIFDTCNTVSKALEATLISFAQNKIDS 122

QY 122 LPIOE--DYSNYISRVAVIGPDNSESVMVTYANFLSLFLLPQITYSATISDELDRKVRPA 179
DB 123 LNLDFCNCSEHIEPTTAVVAGTGCVSTAAANLGLYITIPVSTASSRLSNKQFSS 182
QY 180 LRLTPSADHVEAMVQMLHFRNMWIIVLVSDTYGRDNGQLGERRARRDICIAFOET 239
DB 183 FLRTIPNDHQATNAADIIEYFRMWVGTIAADDYGRGJIEKPEEABEERDICI DFSEL 242
QY 240 LPTLOPNQNMNSEERQRLVTVDKLQOSTARVVVFSGDLTLVAFNEVLRONFGAIVI 299
DB 243 I-----SQYSDDEEIQHVEVI---QNSTAKYIVFSSGPDLEPIKEIVRNITGKIWL 294
QY 300 ASESWAIDPVLANLTELGLTFLGITITQSPPIPGSEF-----REMBQ 344
DB 295 ASEAWASSLLAMPQYFHVVGITGFALKAQIPEGREFLKKVHPRKSVHNGFAKPEWE 354
QY 345 A-----GPPLSRTSOSYTCNQCEDNCNLNTLSFNTILRLSGE----- 382
DB 355 TFNCHLQAGAGPLPVDFTFLRGH--EESGDYFNSNSTAFRL--CTGDENISVETPYID 410
QY 383 ---RVVYSVSAVVAVALHSLIGC-----DKSTCKRVVYPMQLBEIKVNFT 430
DB 411 YTHLRISYVYLVAVSIHALQDIYTCPLGRGLFTNGSCADIKVEAMQVLKHLRLNFT 470
QY 431 -LHDQIFPDQGVALHLEIVQWQMDRSQNP--FQSAVSYPL-----QRLKXIODIS 482
DB 471 NMGEQVTFDECGDLVGVYSIIMNHLSPEDSIVFKEY-GYNNVAKKGERLFTNEKIL 529
QY 483 WHTVNTIPMSMCSKRCQSGOKKPV-GIHVCFECIDCLPQTFNLHTEDEYCOACPNN 541
DB 530 WSGFRSEVPFNSCRDCLAGTRKGIIEBEPCCCEFCVCPGEYDEI--DASACKCDD 588
QY 542 EMSYQSETSCFRQLVLEHHEAPLIAVALLAAGLSTLAILVIFWHPQTPYRSAGG 601
DB 589 FMSNENHTSCIAKEIEFMTPEPGIALTLPAVIGIFLTAFLVGFIFKFRTPVIKATNR 648
QY 602 PMCFMLTLLLVANVNVVYGPVPRVTCRLQALFPLCFITICSCIVRSFOICAF-- 659
DB 649 ELSTLLFLSLCCFSSSLFFIGEPQDWCRLRQAPFISFLVCLISCLIVKTNRLVLEVA 708
QY 660 KMASRFPAYSVWRYOGPVYMAFIVLKNVIVYIGLAPQSHPRTPDPDKITIVSC 719
DB 709 KIPISFHKH--WGNLNLQFLVFLCTFMQIVICIMLYTPAPSSYRQOELEDEIIFITC 765
QY 720 NPNYRNSLLENTSLDLLISVGFSPAYVNGKELPTNNEAKITLSMTFTTSSVSLCTFM 779
DB 766 HEGSIMALGFLIGYTCILAAICFPFAFKSRKLPEFNFAKFTFSMLFIYMWIS--FI 822
QY 780 SAYSGVLTIVDLYVTNLALISLG---YFGPKCYILFYPERNT 822
DB 823 PAYASTYGFVS-AVEVIAIILAAFGLLACIFFNKITYILFKPSRNT 868

RESULT 9
US-08-353-784-7
Sequence 7, Application US/08353784
Patent No. 6011068
GENERAL INFORMATION:
APPLICANT: Edward F. Nemeth, Edward M.
APPLICANT: Brown, Steven C. Hebert,
APPLICANT: Bradford C. Van Wagonen, Manuel
APPLICANT: F. Balandrin, Forrest H. Fuller,
APPLICANT: Eric G. Delmar, and Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESS: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles

FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Douglas C. Murdock
REGISTRATION NUMBER: 37,549
REFERENCE/DOCKET NUMBER: 213/007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-484-719B-7

Query Match 24.6%; Score 1094; DB 3; Length 1078;
Best Local Similarity 31.5%; Pred. No. 8,4e-95;
Matches 279; Conservative 160; Mismatches 342; Indels 106; Gaps 26;

16 WYLAFAENSDPYLP-----GDVLLGLFELHNMGIYVNLQVMEKYEYKYG 68
8 WYLAALWHTSAVYGDQRAQKKDITLLGLFPIHF--GVAKD--QDLKSPRESVECR 62
69 YNL-----MOAMRPAVEEINNDSLLPGVLGYEIVDVC-VISNNVQPLVFLA-HEBVL 121
63 YNFRGFRMLQAMITFAIEEINSSPALPNTLTGRIPTDCTNYSKALEATLSVAQNKIDS 122
122 LPIDQ--DYSNYISRVAVIGPDNSESVTANFLSLFLLPQITTSALSDELDRKVRPFA 179
123 LNLDFCNCSEHIPPSTIAVVGATGSGSTAVANLGLFYITPQVSASSRLLSNKQFQS 182
180 LIRTPSADHVEANVQMLHFRNMWITVLSSTYGRDNGQLGERVARIDICIAFOET 239
183 FLRTIPNDEHQATPAADIIEYFRMWVGTIADDDYGRGAIKFEAEERDIDICIDFSEL 242
240 LPTLPQNMNTSEERQRLTIVDKLQOSTARVVVPSPLTLVHFNEVLRNFTGAWVI 299
243 I-----SQYSDDEEIOHVEVI--QNSTAKYIVVSSSPDEPLIKETIVRNITGKJWL 294
300 ASESWAIDPVNLHNLTELGLTGLITTSQVPIPGFSEF-----REWGPO 344
295 ASEAWASSLLAMPQYFHVVGCTIGFALKAGIIPGRREFLKVKHPRKSVHNGFAKEFME 354
345 A-----GPPUSRTSQSYTCNOECDNCLNATLSFNTILRLSGE----- 382
355 TFNCHLOBAKGPLVDVDFLRGH--EESGDRFSNSSTAFRPL--CTGDEMISVETPYID 410
363 -----RVVSVYSAVVAVALHSLGCG-----DKSCTRVVYVPMOLLEIKUNAT 430
411 YTHLISINIVAVYVSIHALQDIYTCPLGRGLFTNGSCADIKKVEAMQVHLHLHNT 470
431 -LLDHIQIFDDPGDVALHLEIVQWQMDRSQNP--FQSVASYVPL-----QROLKNIODIS 482
471 NNMGGQVTFDECGDLVGNYSIINMHLSPEDGSIVRKEV--GYINNVAKKGERLFINBEKTL 529

483 WHTVNTIPMSGSKRCSGOKKKV--GIHVCCFECIDCLPOTPLNHTDEYECQACPN 541
530 MSGFRVFPFNCSDRCDLACGRKGIIBEPPTCCFCEVBCPGEYDEI--DASACKCD 588
542 EMSYQSTSCFRQQLVLEHAPTAVALAAGFSLTALVIFEMHFPQTPYRSAG 601
589 FWSNMHTSCIAKEIEFLSWTEPFQIALTLFAVIGIFLTAAGVIFRNPPIKATNR 648
602 FMCFMLTLLVAVNVVYVGPVSTCLRQALFPCFPTICISCIIVRSQIYCAF-- 659
649 ELSTYLLFSLCCRSSSLFFIAGEPDWTCRLRQAPFGLISFVLCISCLIVKTRVLV 708
660 KMASRFPAYSVWVRVYQPYVSMFITYLKNVIVIGKLARPOSIPRTDPPDKITVISC 719
709 KIPFSFHK--WGLNMQFLVFLCTMJOIVCYIMLYTAPSSYRNQLEDELIIFITC 765
720 NPNYRNSLLFNTSIDLLSVGFSPAYVSKELPTVYNAKFTLSMTFTSSVSLCTFM 779
766 HEGSLMALGFLIGVTCLLAALCFEAFKSRKLPENFNAKFTTFMILFIWIS---FI 822
780 SAYSGVLTIVDLYTVNLALISIG---YFGPKYMLFYPPENT 822
823 PAYASTYGRFVS-AVEVTAIILAASFGILACIFFNKIYIILFRPSHNT 868

RESULT 11
US-08-484-159-7
Sequence 7, Application US/08484159
Patent No. 6313146
GENERAL INFORMATION:
APPLICANT: Bradford C. Van Magenen
APPLICANT: Manuel F. Balandrin
APPLICANT: Eric G. Del Mar
APPLICANT: Edward F. Nemeth
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,159
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 9
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992

APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 214/101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-484-159-7

Query Match 24.6%; Score 1094; DB 4; Length 1078;

Best Local Similarity 31.5%; Pred. No. 8,4e-95;
Matches 279; Conservative 160; Mismatches 342; Indels 106; Gaps 26;

16 WLAEPANSDPYLP-----GDYLLGLFSLHANMKGIYHNLQVPMCKEYEVKIG 68
8 WVLALHTHTSAYGDDQAKKGDIIIGLFPPIHF---GVAKD--QDLKSRPSEVEIR 62
69 YNL-----MAMRFAVEEINDDSLPGVLLGYEIVDVC-YISNNQPVLYLA-HEENL 121
63 YNFRGFWLQAMIFAEIEINSPALLPVLTLGYRIFDCTNVSFKALEATLSFVQNKIDS 122
122 LPICE--DYSNYISRVVAVGPNSESVMTANFLSLFLLPQITYSASIDELRDVRFPA 179
123 LNLDFCNCSEHIEPTIAVAGATGCVSTAVANLGLFYIPVSTYASSRLISNCKPKS 182
180 LRLTPSADHVEANVQMLHFRNMWIIYLVSSDTYGRDNGQLGERVARRDICTAFOET 239
183 FLRTIPNEHQATAMADIEYFRMNVGTIAADDYGRGIEKFEAEAEEDICIDFSEL 242
240 LPTLPONQMTSEERQRLVTVDKLQOSTARVVVPSDLTYHFEENEVLRONFGAVWI 299
243 I-----SQYSDDEEIQHVEVY---QNSTAKIVVFSGGPDLEPLIKEIVRNITGKTWL 294
300 ASESWAIDPVNLNLDELGLGTLGITIOSVPIPGSEF-----REMGPO 344
295 ASEAWASSLIAMPQYFHVVGSTIGFALKAQIPGFREBLKVKVRKSVHNGFAKEFWES 354
345 A-----GPPPLSRTSQSYTCNOECNCLNATLSFTILRLSGE----- 382
355 TFNCHLGAGKGPLVDTFRLGH--EESGDRFSNSTAFRLP--CTGDENTSSVETPYID 410
383 ----RVVYSVSYAVVAHALHSLIGC-----DKSTCKRVVYPMQLBEIKVNF 430
411 YTHLRISYVAVVAVSIHALQDIYTCPLGRGLFTNGSCADIKKVEAMQVKKHMLNFT 470
431 -LLHQIEPDDQGVALLLEIVQWMDRSQNP--FQSVASYRPL-----OQLKQIDIS 482
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530 WSGREVEYFNSCSDCLAGTRKGIIEBEPCCFCEVCBPDEYEDFI-DASACKKCPDD 588
589 FWSNENHTSCIAKEIEFISWTEPFGIALTLFAVLGIFLTAFLGVIFIKFRTPPIVAKNR 648
542 EMSYQSESTCFKRLQVLEMHAPTIIVALLAALGFLSTLALIVFMHFQTPPIVRSAGG 601
602 PMCFMLLLLVAVMVVVPVYVCPKVSCTCROALFPLCFTTICSCIVRSPQIYCAR-- 659
649 ELSYLLSLSLCCFSSSLFPIGEPQDMTCRLRQPAFGISFVLCISCIIVKTNRLVLEBA 708
660 KMASRFPAYSVWVRYOGPVYVMAFETVLKMWIVIGMLARPOSHRTPDPDKITIVSC 719

709 KIPTSFARK---WMGLNLOFLVFLCTFMOIVICVIMLYTAPSSSYANOELIEBIIPTC 765
720 NPNRNSLLENTSIDLLSVGPFSAWKELPNVEAKFTLSMFFYFTSSSLCTFM 779
766 HEGSLMALGFIYGTCTLLAICFFFAKSRKLPENFEAKFTFSMLIFVWIS---FI 822
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823 PAVASTYKGFVS-AVEVIALILASFGLLACIFPKXIIILFKPSRNT 866

RESULT 12
US-09-162-021B-2

Sequence 2, Application US/09162021B

Patent No. 6337391

GENERAL INFORMATION:

APPLICANT: H. William Harris

APPLICANT: Edward M. Brown

APPLICANT: Steven C. Hebert

TITLE OF INVENTION: Polyclonal-Sensing Receptor in Aquatic

TITLE OF INVENTION: Species and Methods of Use Thereof

FILE REFERENCE: 2856.1001-007

CURRENT FILING DATE: 1998-09-28

PRIOR APPLICATION NUMBER: PCT/US97/05031

PRIOR FILING DATE: 1997-03-27

PRIOR APPLICATION NUMBER: 08/622,738

PRIOR FILING DATE: 1996-03-27

NUMBER OF SEQ ID NOS: 19

SOFTWARE: FastSeq for Windows Version 4.0

SEQ ID NO 2

LENGTH: 1027

TYPE: PRT

ORGANISM: aequi

US-09-162-021B-2

Query Match 24.6%; Score 1092.5; DB 4; Length 1027;

Best Local Similarity 31.0%; Pred. No. 1.1e-94;
Matches 277; Conservative 162; Mismatches 360; Indels 95; Gaps 25;

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2 AQLHCQLLFLGFTLLQ--SYVSGYGPQRAQKKGDIILGLFPIHFGVAAKDDOLKSRPE 60
56 VPMCKEYEVKIGNLMQAMRFAVEEINDDSLPGVLLGYEIVDVC-YISNNQPVLYF 114
61 ATKCIKRYNFR--GRWLQAMIFAEIEINSMFTFLPNTLIGRIFDCTNVSFKALEATLSF 118
115 LA-HEENLPIQE--DYSNYISRVVAVIGPNSESVMTANFLSLFLLPQITYSASIDEL 171
119 VQNKIDSLNDEHCNCSDDHPSITIAVAGATGSIATAVANLGLFYIPVSTYASSRL 178
172 RDKVRPALLRTPSADHVEANVQMLHFRNMWIIYLVSSDTYGRDNGQLGERVARRD 231
179 SNKVEYKAPLTIIPNDQATAMAEITEHFQWVVGTLAADDDYGRGIDKFEAEAVKRD 238
232 ICTAFOETLPLQONQMTSEERQRLVTVDKLQOSTARVVVPSDLTYHFEENEVLRQ 291
239 ICIDFSEMI-----SQYTT---QKQLEFIADVIONSSAKVIVFNSGPDLEPLIQEIVR 290
292 NFGAVWIASESMAIDVNLNLDELGLGTLGITIOSVPIPGSEFPE----- 340
291 NITRIVLASEAMASSSLIAKPEYFHVVGSTIGFALPAAGIIPGFNKLKVEHPSRSSDNG 350
341 ----W-----GPOA--GPPPLSRTSQSYTCNOECNCLNAT 370
351 FVKEFMEETFNCFYTEXTLQNLKSKVPSHGPAQGGGSKAGNSRFRALRHPCTGEBNIT 410
371 LSFNTIIRLGERRVVYSVAVVAHALHSLIGCDST-----CTK--RVVYPMQLLE 422
411 SVETPYLDYTHLRISYVAVVAVSIHALQDIHSCCKPQGTGIFANGSCADIKKVEAMQV 470

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GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: May 11, 2004, 15:27:42 ; Search time 40.6604 Seconds
(without alignments)
5720.583 Million cell updates/sec

Title: US-09-927-315-9

Perfect score: 4443

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Scoring table:

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Gapop 10.0 , Gapext 0.5

Searched: 1140673 seqs, 27756755 residues

Total number of hits satisfying chosen parameters: 1140673

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	4443	100.0	838	14	US-10-190-417-9
3	4392.5	98.9	839	9	US-09-897-427A-4
4	4392.5	98.9	839	14	US-10-035-045-21
5	4392.5	98.9	839	15	US-10-179-373-6
6	4372.5	98.4	839	14	US-10-246-785-4
7	3514	79.1	669	13	US-10-124-598-7
8	3514	79.1	669	14	US-10-096-144-7
9	3514	79.1	669	14	US-10-225-567A-683
10	3203.5	72.1	843	13	US-09-927-315-7
11	3203.5	72.1	843	13	US-10-134-598-1
12	3203.5	72.1	843	14	US-10-096-144-1
13	3203.5	72.1	843	14	US-10-246-785-6
14	3203.5	72.1	843	14	US-10-190-417-7
15	3203.5	72.1	843	15	US-10-179-373-17

16	3203.5	72.1	843	15	US-10-436-715-38	Sequence 38, Appl
17	3203.5	72.1	843	15	US-10-436-715-70	Sequence 70, Appl
18	3151.5	70.9	843	10	US-09-927-315-8	Sequence 8, Appl
19	3151.5	70.9	843	13	US-10-124-598-2	Sequence 2, Appl
20	3151.5	70.9	843	14	US-10-096-144-2	Sequence 2, Appl
21	3151.5	70.9	843	14	US-10-190-417-8	Sequence 8, Appl
22	2505	56.4	661	14	US-10-246-785-5	Sequence 5, Appl
23	1454	32.7	840	10	US-09-361-652-1	Sequence 1, Appl
24	1454	32.7	840	10	US-09-927-315-1	Sequence 1, Appl
25	1454	32.7	840	14	US-10-246-785-3	Sequence 3, Appl
26	1454	32.7	840	14	US-10-190-417-1	Sequence 1, Appl
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29	1454	32.7	840	15	US-10-436-715-39	Sequence 39, Appl
30	1454	32.7	840	15	US-10-436-715-69	Sequence 69, Appl
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32	1440	32.4	842	15	US-10-436-715-67	Sequence 67, Appl
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36	1434	32.3	842	14	US-10-190-417-2	Sequence 2, Appl
37	1431	32.2	842	15	US-10-436-715-68	Sequence 68, Appl
38	1354	30.5	841	9	US-09-819-946-2	Sequence 2, Appl
39	1354	30.5	841	9	US-09-897-427A-2	Sequence 2, Appl
40	1354	30.5	841	10	US-09-799-629-17	Sequence 17, Appl
41	1354	30.5	841	14	US-10-035-045-17	Sequence 17, Appl
42	1354	30.5	841	14	US-10-190-417-27	Sequence 27, Appl
43	1354	30.5	841	15	US-10-179-373-5	Sequence 5, Appl
44	1353	30.5	841	12	US-10-188-186-2	Sequence 2, Appl
45	1336.5	30.1	840	14	US-10-190-417-3	Sequence 3, Appl

ALIGNMENTS

RESULT 1
US-09-927-315-9
Sequence 9, Application US/09927315
Publication No. US20030040045A1
GENERAL INFORMATION:
APPLICANT: Zuker, Charles S.
APPLICANT: Ryda, Nicholas J.P.
APPLICANT: Nelson, Greg
APPLICANT: Hoon, Mark A.
APPLICANT: Chandrasekhar, Jayaram
APPLICANT: Zhang, Yiteng
APPLICANT: The Regents of the University of California
APPLICANT: The Government of the United States of America
APPLICANT: as represented by the Secretary of the
TITLE OF INVENTION: Mammalian Sweet Taste Receptors
FILE REFERENCE: 02307E-120110US
CURRENT APPLICATION NUMBER: US/09/927,315
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: US 60/302,898
PRIOR FILING DATE: 2001-07-03
NUMBER OF SEQ ID NOS: 25
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 9
LENGTH: 838
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: human TIR2 sweet taste receptor
US-09-927-315-9

Query Match 100.0%; Score 4443; DB 10; Length 838;

Best local Similarity 100.0%; Pred. NO. 0; Mismatches 0; Indels 0; Gaps 0;

Matches 838; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Db 1 MGPRAKTICSLFLLMWLAEPNSDFLLPQDYLGGIFSLHAKMKGIIVHLNFIQVPMCK 60

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RESULT 2
US-10-190-417-9
; Sequence 9, Application US/10190417
; Publication No. US20030166137A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Ryda, Nicholas J. P.
; APPLICANT: Chandrasekar, Jayaram
; APPLICANT: Hoon, Mark A.
; APPLICANT: Nelson, Greg
; APPLICANT: Zhang, Yifeng
; APPLICANT: The Regents of the University of California
; APPLICANT: The Government of the United States of America
; APPLICANT: as represented by the Secretary of the
; APPLICANT: Department of Health and Human Services
; TITLE OF INVENTION: Mammalian Sweet and Amino Acid Heterodimeric Taste
; TITLE OF INVENTION: Receptors
; FILE REFERENCE: 02307E-120130US
; CURRENT APPLICATION NUMBER: US/10-190, 417
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; CURRENT FILING DATE: 2002-11-14
; PRIOR APPLICATION NUMBER: US 60/302, 898
; PRIOR FILING DATE: 2001-07-03
; PRIOR APPLICATION NUMBER: US 09/927, 315
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: US 60/358, 925
; PRIOR FILING DATE: 2002-02-22
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 838
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: human T1R2
US-10-190-417-9

Query Match 100.0%; Score 4443; DB 14; Length 838;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 838; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 3
US-09-897-427A-4
; Sequence 4, Application US/09897427A
; Patent No. US2002016042A1
; GENERAL INFORMATION:
; APPLICANT: ADLER, JON ELLIOT
; APPLICANT: LI, XIADONG
; APPLICANT: STAZEMSKI, LENA
; APPLICANT: XU, HONG
; APPLICANT: EHEVERRI, FERNANDO
; TITLE OF INVENTION: T1R HETERO-OLIGOMERIC TASTE RECEPTORS
; FILE REFERENCE: 078003-0282558
; CURRENT APPLICATION NUMBER: US/09/897,427A
; CURRENT FILING DATE: 2001-07-03
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 839
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-897-427A-4

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Dp	601	GPMSFLMTLLIVAAVVVYVYGGPRVSTCLCRQALPYLCFTTICISCIANRSFOIVCAFK	660	

Db	601	GMCFMLTLTLVAVMVPVYVGPVKVSTCLGRQALPFCFTICISICIAVRSQIVCAFK	660
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Qy	720	NPVYNSLLFNTSLDILLSVGSEFPAWYMKELPTNNNAKFTILSMTPVFTSSVSLCTFM	779
Db	721	NPVYNSLLFNTSLDILLSVGSEFPAWYMKELPTNNNAKFTILSMTPVFTSSVSLCTFM	780
Qy	780	SAYSGVLVTIYDILLVTLVNLALISIGYFQPKCYMILFPERNTPAYFNSMIGQYTRMD	838
Db	781	SAYSGVLVTIYDILLVTLVNLALISIGYFQPKCYMILFPERNTPAYFNSMIGQYTRMD	839

```

RESULT 4
US-10-035-045-21
; Sequence 21, Application US/10035045
; Publication No. US20030054448A1
; GENERAL INFORMATION:
; APPLICANT: ADLER, JON ELLIOT
; APPLICANT: LI, XIAODONG
; APPLICANT: STASZEWSKI, LENA
; APPLICANT: O'CONNELL, SHAWN
; APPLICANT: ZOZULIYA, SERGEY
; TITLE OF INVENTION: T1R TASTE RECEPTORS AND GENES ENCODING SAME
; FILE REFERENCE: 078003-0280681
; CURRENT APPLICATION NUMBER: US/10/035,045
; CURRENT FILING DATE: 2002-01-03
; PRIOR APPLICATION NUMBER: 60/259,227
; PRIOR FILING DATE: 2001-01-03
; PRIOR APPLICATION NUMBER: 60/284,547
; PRIOR FILING DATE: 2001-04-19
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 21
; LENGTH: 839
; TYPE: prt
; ORGANISM: Homo sapiens
US-10-035-045-21

```

Query Match	Similarity	98.9%	Score 4392.5	DB 14	Length 839
Best Local	Similarity	99.3%	Pred. No. 0		
Matches	833	Conservative	0	Mismatches	5
				Indels	1
				Gaps	1
QY	1	MGPRAKTICSLEFLLMTAEPRAENSDFYLPDYLGGFLSHLANKGIYHNLQVPMCK	60		
DB	1	MGPRAKTICSLEFLLMTAEPRAENSDFYLPDYLGGFLSHLANKGIYHNLQVPMCK	60		
QY	61	EYEVAVIGNLMQARFAVEEINNDSLLPGVLGVEIYDVCIYNNVQVYFLAHEDN	120		
DB	61	EYEVAVIGNLMQARFAVEEINNDSLLPGVLGVEIYDVCIYNNVQVYFLAHEDN	120		
QY	121	LIPLOEDYSNYISRVAVYIGPDNSESVMTVANFLSLPILPQITYSAISDELDRKVRPAL	180		
DB	121	LIPLOEDYSNYISRVAVYIGPDNSESVMTVANFLSLPILPQITYSAISDELDRKVRPAL	180		
QY	181	LRTTESADHHVAMVQOLMHFRPMNNIIVVSSPTYRDNGQLLGEAVARDDICAFQETL	240		
DB	181	LRTTESADHHVAMVQOLMHFRPMNNIIVVSSPTYRDNGQLLGEAVARDDICAFQETL	240		
QY	241	PLLOPNQNTSEEROLVTIYDKLOOSTARVVVYSPDLTYHFNVEVLRQNTFGAVWIA	300		
DB	241	PLLOPNQNTSEEROLVTIYDKLOOSTARVVVYSPDLTYHFNVEVLRQNTFGAVWIA	300		
QY	301	SESMAIDPVLNHLTELGHGTFLGTTIGSVPIPGSEFEFREMGOAAPPPLSRFSQSQYTCN	360		
DB	301	SESMAIDPVLNHLTELGHGTFLGTTIGSVPIPGSEFEFREMGOAAPPPLSRFSQSQYTCN	360		
QY	361	QECNDCLNATLSFNITLRLSGRRVYVYVSAYAAVAHLSLLGCKSKTCKRVVYPMOL	420		
DB	361	QECNDCLNATLSFNITLRLSGRRVYVYVSAYAAVAHLSLLGCKSKTCKRVVYPMOL	420		

Qy 421 LEEIWKVNFLLDHOIFPDPOGDVALHLEIVQMOMDRSQNPFQSVASYYPLOQLKNIQD 480
Db 421 LEEIWKVNFLLDHOIFPDPOGDVALHLEIVQMOMDRSQNPFQSVASYYPLOQLKNIQD 480
Qy 481 ISMHTVNNITIPMSKSGKQSGQKKKPVGIVHCCFECIDCLPGTFLNTEDEYECQACP 540
Db 481 ISMHTVNNITIPMSKSGKQSGQKKKPVGIVHCCFECIDCLPGTFLNTEDEYECQACP 540
Qy 541 NEMSYQSETSCFKROLVLEHMEAPTIYVALLAALGFLSTLAIIVFMRHFOPIVRSG 600
Db 541 NEMSYQSETSCFKROLVLEHMEAPTIYVALLAALGFLSTLAIIVFMRHFOPIVRSG 600
Qy 601 GPMCFMLTLTLLVAVVYVPPVGPVKSTCLCQALPFLCFTICISCIASVSPQIVCAF 660
Db 601 GPMCFMLTLTLLVAVVYVPPVGPVKSTCLCQALPFLCFTICISCIASVSPQIVCAF 660
Qy 661 MASRFPAYSYWRYQGPVSMAFITVLKXVIVVIGMLARPOS-HPRTPDDPKITIVSC 719
Db 661 MASRFPAYSYWRYQGPVSMAFITVLKXVIVVIGMLARPOS-HPRTPDDPKITIVSC 720
Qy 720 NPNYRNSLLENTSLDLTSLVGFSPAYMGKELPTVYNEAKFITLSMTFFTSVSLCTFM 779
Db 720 NPNYRNSLLENTSLDLTSLVGFSPAYMGKELPTVYNEAKFITLSMTFFTSVSLCTFM 780
Qy 780 SAYSGVLVTIVDLVTVLNLALISLGYFGPKCYMILFYPERNTPAYFNSMIQGYTMRD 838
Db 781 SAYSGVLVTIVDLVTVLNLALISLGYFGPKCYMILFYPERNTPAYFNSMIQGYTMRD 839

RESULT 5

US-10-179-373-6

Sequence 6, Application US/10179373

Publication No. US20030232407A1

GENERAL INFORMATION:

APPLICANT: ZOLLER, MARK

APPLICANT: LI, XIAODONG

APPLICANT: STASZEWSKI, LENA

APPLICANT: O'CONNELL, SHAWN

APPLICANT: ZOZULYA, SERGEY

APPLICANT: ADLER, JON

APPLICANT: XU, HONG

TITLE OF INVENTION: T1R HETERO-OLIGOMERIC TASTE RECEPTORS AND CELL LINES

TITLE OF INVENTION: THAT EXPRESS SAID RECEPTORS AND USE THEREOF FOR

TITLE OF INVENTION: IDENTIFICATION OF TASTE COMPOUNDS

FILE REFERENCE: 078003-0291566

CURRENT APPLICATION NUMBER: US/10/179.373

PRIOR FILING DATE: 2002-06-26

PRIOR APPLICATION NUMBER: 60/300,434

PRIOR FILING DATE: 2001-06-26

PRIOR APPLICATION NUMBER: 60/304,749

PRIOR FILING DATE: 2001-07-13

PRIOR APPLICATION NUMBER: 60/310,493

PRIOR FILING DATE: 2001-08-08

PRIOR APPLICATION NUMBER: 60/331,771

PRIOR FILING DATE: 2001-11-21

PRIOR APPLICATION NUMBER: 60/339,472

PRIOR FILING DATE: 2001-12-14

PRIOR APPLICATION NUMBER: 60/372,090

PRIOR FILING DATE: 2002-04-15

PRIOR APPLICATION NUMBER: 60/374,143

PRIOR FILING DATE: 2002-04-22

NUMBER OF SEQ ID NOS: 19

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 6

LENGTH: 839

TYPE: PRT

ORGANISM: Homo sapiens

US-10-179-373-6

Query Match 98.9%; Score 4392.5; DB 15; Length 839;
Best Local Similarity 99.3%; Pred. No. 0;
Matches 833; Conservative 0; Mismatches 5; Indels 1; Gaps 1;

Qy 1 MGPRAKTICSLFLLWVLAEPANSDPYLPQDYLGLGFLSHANKMGIHNLNFQVEMCK 60
Db 1 MGPRAKTICSLFLLWVLAEPANSDPYLPQDYLGLGFLSHANKMGIHNLNFQVEMCK 60
Qy 61 EYEVKIVGYMLQMPRAVEEINNDSLLPQVLGYEIVDVCY1SNVQPVLYFLAEDN 120
Db 61 EYEVKIVGYMLQMPRAVEEINNDSLLPQVLGYEIVDVCY1SNVQPVLYFLAEDN 120
Qy 121 LLPQEDYSNYSISRVVAVIGDSESVMTANFLSLPQITYSALSDELKDYRFPAL 180
Db 121 LLPQEDYSNYSISRVVAVIGDSESVMTANFLSLPQITYSALSDELKDYRFPAL 180
Qy 181 LRTTSPADHVEAVQMLHFRMNMIIVLVSDTYGRDNGQLGERVARRDICIAPQETL 240
Db 181 LRTTSPADHVEAVQMLHFRMNMIIVLVSDTYGRDNGQLGERVARRDICIAPQETL 240
Qy 241 PTLQPNQNMTESEERQLVTVDKLQOSTARVVVSPDLTYHPFNEVLBQNTGAVVIA 300
Db 241 PTLQPNQNMTESEERQLVTVDKLQOSTARVVVSPDLTYHPFNEVLBQNTGAVVIA 300
Qy 301 SESWADPVVNLHNLTELHGLTFLGITIQSVPIRGSFSEFRMGQAGPPLSRTSQSTCN 360
Db 301 SESWADPVVNLHNLTELHGLTFLGITIQSVPIRGSFSEFRMGQAGPPLSRTSQSTCN 360
Qy 361 QECNCLNATLSFTTILRLSGERYVYSVAVYAAHALLSLGCDKSTCTKRVVYPMQL 420
Db 361 QECNCLNATLSFTTILRLSGERYVYSVAVYAAHALLSLGCDKSTCTKRVVYPMQL 420
Qy 421 LEEIWKVNFLLDHOIFPDPOGDVALHLEIVQMOMDRSQNPFQSVASYYPLOQLKNIQD 480
Db 421 LEEIWKVNFLLDHOIFPDPOGDVALHLEIVQMOMDRSQNPFQSVASYYPLOQLKNIQD 480
Qy 481 ISMHTVNNITIPMSKSGKQSGQKKKPVGIVHCCFECIDCLPGTFLNTEDEYECQACP 540
Db 481 ISMHTVNNITIPMSKSGKQSGQKKKPVGIVHCCFECIDCLPGTFLNTEDEYECQACP 540
Qy 541 NEMSYQSETSCFKROLVLEHMEAPTIYVALLAALGFLSTLAIIVFMRHFOPIVRSG 600
Db 541 NEMSYQSETSCFKROLVLEHMEAPTIYVALLAALGFLSTLAIIVFMRHFOPIVRSG 600
Qy 601 GPMCFMLTLTLLVAVVYVPPVGPVKSTCLCQALPFLCFTICISCIASVSPQIVCAF 660
Db 601 GPMCFMLTLTLLVAVVYVPPVGPVKSTCLCQALPFLCFTICISCIASVSPQIVCAF 660
Qy 661 MASRFPAYSYWRYQGPVSMAFITVLKXVIVVIGMLARPOS-HPRTPDDPKITIVSC 719
Db 661 MASRFPAYSYWRYQGPVSMAFITVLKXVIVVIGMLARPOS-HPRTPDDPKITIVSC 720
Qy 720 NPNYRNSLLENTSLDLTSLVGFSPAYMGKELPTVYNEAKFITLSMTFFTSVSLCTFM 779
Db 720 NPNYRNSLLENTSLDLTSLVGFSPAYMGKELPTVYNEAKFITLSMTFFTSVSLCTFM 780
Qy 780 SAYSGVLVTIVDLVTVLNLALISLGYFGPKCYMILFYPERNTPAYFNSMIQGYTMRD 838
Db 781 SAYSGVLVTIVDLVTVLNLALISLGYFGPKCYMILFYPERNTPAYFNSMIQGYTMRD 839

RESULT 6

US-10-246-785-4

Sequence 4, Application US/10246785

Publication No. US20030148448A1

GENERAL INFORMATION:

APPLICANT: IRM, LLC

APPLICANT: The Scripps Research Institute

APPLICANT: Liao, Jiayu

APPLICANT: Sheng, Ding

APPLICANT: Schultz, Peter G

TITLE OF INVENTION: Sweet Taste Receptors

FILE REFERENCE: 36-002810US/PC

CURRENT APPLICATION NUMBER: US/10/246.785

CURRENT FILING DATE: 2002-12-09

PRIOR APPLICATION NUMBER: US 60/323,450

PRIOR FILING DATE: 2001-09-18
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 839
TYPE: PRT
ORGANISM: Homo sapiens
US-10-246-795-4

Query Match 98.4%; Score 4371.5; DB 14; Length 839;
Best Local Similarity 98.7%; Pred. No. 0;
Matches 828; Conservative 2; Mismatches 8; Indels 1; Gaps 1;

1 MPPRAKTIQSFLFVLVLAEPANSPFYLPQYLLGGLSLHANKKGIHVLPLQVPMCK 60
1 MPPRAKTIQSFLFVLVLAEPANSPFYLPQYLLGGLSLHANKKGIHVLPLQVPMCK 60
61 EYEVKVIQYINLMQAMFAVEEINNDSSLLPGVLLAGEIVDVCYISNNQPVLYFLAHEDN 120
61 EYEVKVIQYINLMQAMFAVEEINNDSSLLPGVLLAGEIVDVCYISNNQPVLYFLAHEDN 120
121 LIPIDEDYSNYSIRVAVAVIGPNSESVMTVANFLSLFLPQITTSASISDELBDKXRFPAL 180
121 LIPIDEDYSNYSIRVAVAVIGPNSESVMTVANFLSLFLPQITTSASISDELBDKXRFPAL 180
181 LATTSSADHVEAMQMLHFRNMNTIYVSSDTYGRDNGQLGGERVARDICIAFOETL 240
181 LATTSSADHVEAMQMLHFRNMNTIYVSSDTYGRDNGQLGGERVARDICIAFOETL 240
241 PTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTLVHFNEVLKQNFQAVMIA 300
241 PTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTLVHFNEVLKQNFQAVMIA 300
301 SESMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEWGPQAGPPPLSRTSOSYTCN 360
301 SESMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEWGPQAGPPPLSRTSOSYTCN 360
361 OECDNCNANTLSFNTILRISGERVYVSVYSAVAHAHLSLGCDSKCTCRVYVPMOL 420
361 OECDNCNANTLSFNTILRISGERVYVSVYSAVAHAHLSLGCDSKCTCRVYVPMOL 420
421 LEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONPQSVASYPLOROLKNIOD 480
421 LEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONPQSVASYPLOROLKNIOD 480
481 ISMHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCLPOTFLNHTEDYECCACPN 540
481 ISMHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCLPOTFLNHTEDYECCACPN 540
541 NEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTALIVYFMHFGPTIVRSAG 600
541 NEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTALIVYFMHFGPTIVRSAG 600
601 GPMCLMLTLVAVAVVYVGPVKVSTCLCRQALFPLCTITICISCIASVRSFOIVCAFK 660
601 GPMCLMLTLVAVAVVYVGPVKVSTCLCRQALFPLCTITICISCIASVRSFOIVCAFK 660
661 MASRPRAVSVMVYOGPYVMAFITVLMKVIIVIGMLATGSLPTRIDDDPKITIVSC 719
661 MASRPRAVSVMVYOGPYVMAFITVLMKVIIVIGMLATGSLPTRIDDDPKITIVSC 719
720 NNNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTLTSMTFFTSVSLCTFM 779
720 NNNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTLTSMTFFTSVSLCTFM 779
780 SAYSGVLVTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 838
780 SAYSGVLVTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 838
839 SAYSGLVTVLTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 839

RESULT 7
US-10-124-598-7
Sequence 7, Application US/10124598

Publication No. US20020119526A1
GENERAL INFORMATION:
APPLICANT: Zuker, Charles S.
APPLICANT: Adler, Jon Elliot
APPLICANT: Lindemeier, Uergeren
TITLE OF INVENTION: The Regents of the University of California
TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
FILE REFERENCE: 02307E-088720US
CURRENT APPLICATION NUMBER: US/10/124,598
PRIORITY FILING DATE: 2002-04-16
PRIORITY FILING DATE: EARLIER APPLICATION NUMBER: US/09/361,631
PRIORITY FILING DATE: EARLIER FILING DATE: 1999-07-27
PRIORITY FILING DATE: EARLIER APPLICATION NUMBER: US 60/112,747
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7
LENGTH: 669
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: human G-protein coupled receptor (GPCR) B4 amino
US-10-124-598-7

Query Match 79.1%; Score 3514; DB 13; Length 669;
Best Local Similarity 98.4%; Pred. No. 0;
Matches 666; Conservative 1; Mismatches 2; Indels 8; Gaps 2;

162 ITYSASIDELBDKXRFPALRTTPSADHVEAMQMLHFRNMNTIYVSSDTYGRDNGQ 221
1 ITYSASIDELBDKXRFPALRTTPSADHVEAMQMLHFRNMNTIYVSSDTYGRDNGQ 60
222 LIGERVARDICIAFOETLPTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTL 281
222 LIGERVARDICIAFOETLPTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTL 281
61 LIGERVARDICIAFOETLPTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTL 120
61 LIGERVARDICIAFOETLPTLQPNQNMNTSEERORLVTIYDKLQOSTARVVVVFSPDLTL 120
282 YHFEVNLKQNFQAVMIASSMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEW 341
282 YHFEVNLKQNFQAVMIASSMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEW 341
121 YHFEVNLKQNFQAVMIASSMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEW 180
121 YHFEVNLKQNFQAVMIASSMAIDPVLANLTELGHGTLGTLTIGSVPIPGSEFEFEW 180
342 GPQAGPPPLSRTSOSYTCNOECDNCNANTLSFNTILRISGERVYVSVYSAVAHAHLS 401
342 GPQAGPPPLSRTSOSYTCNOECDNCNANTLSFNTILRISGERVYVSVYSAVAHAHLS 401
401 GPQAGPPPLSRTSOSYTCNOECDNCNANTLSFNTILRISGERVYVSVYSAVAHAHLS 240
402 LIGCDKSTCRVYVPMOLLEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONP 461
402 LIGCDKSTCRVYVPMOLLEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONP 461
241 LIGCDKSTCRVYVPMOLLEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONP 300
241 LIGCDKSTCRVYVPMOLLEEIKVNFETLLDHOIFPDPOGVALHLEIVOMQDRSONP 300
462 FOSVASVYPLOROLKNIODISMHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCL 521
462 FOSVASVYPLOROLKNIODISMHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCL 521
301 FOSVASVYPLOROLKNIK-TSLHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCL 359
301 FOSVASVYPLOROLKNIK-TSLHTVNTIIPMSCKSKRCQSGQKKKPVGIHVCCEFCIDCL 359
522 PGTELNHTEDYECCACPNEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTL 581
522 PGTELNHTEDYECCACPNEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTL 581
360 PGTELNHTEDYECCACPNEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTL 412
360 PGTELNHTEDYECCACPNEMSVOSSESCFRQOLVLEHMEAPRTIAVALLAAGFLSTL 412
582 AILVIFMHPOTPIVRSAGPMLTLVAVAVVYVGPVKVSTCLCRQALFPLCF 641
582 AILVIFMHPOTPIVRSAGPMLTLVAVAVVYVGPVKVSTCLCRQALFPLCF 641
413 AILVIFMHPOTPIVRSAGPMLTLVAVAVVYVGPVKVSTCLCRQALFPLCF 472
413 AILVIFMHPOTPIVRSAGPMLTLVAVAVVYVGPVKVSTCLCRQALFPLCF 472
642 TITICISCIASVRSFOIVCAFKMASRPRAVSVMVYOGPYVMAFITVLMKVIIVIGMLARP 701
642 TITICISCIASVRSFOIVCAFKMASRPRAVSVMVYOGPYVMAFITVLMKVIIVIGMLARP 701
702 OSHPRTDDBDKITIVSCNPNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTI 761
702 OSHPRTDDBDKITIVSCNPNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTI 761
533 OSHPRTDDBDKITIVSCNPNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTI 592
533 OSHPRTDDBDKITIVSCNPNYNSNLLFNTSLDLLSVGFSFAYMKELPNTYNEAKFTI 592
762 TLSMTFFTSVSLCTFMSASVGLVTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 821
762 TLSMTFFTSVSLCTFMSASVGLVTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 821
593 TLSMTFFTSVSLCTFMSASVGLVTVLTVLTVLNLALASLGFGPKCYMLIFPERBTPAYFNSMIOGYTMRD 652

Oy 822 TPAYNSMIOGYTMRD 838
Db 653 TPAYNSMIOGYTMRD 669

RESULT 8

US-10-096-144-7
; Sequence 7, Application US/10096144
; Publication No. US2003002288A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliot
; APPLICANT: Lindemeier, Juergen
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; TITLE OF INVENTION: Involved in Sensory Transduction
; FILE REFERENCE: 02307E-088720US
; CURRENT APPLICATION NUMBER: US/10/096,144
; PRIOR FILING DATE: 2002-03-08
; PRIOR APPLICATION NUMBER: EARLIER FILING DATE: 1999-07-27
; PRIOR FILING DATE: EARLIER FILING DATE: 1998-07-27
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/095,464
; PRIOR FILING DATE: EARLIER FILING DATE: 1998-07-28
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/112,747
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO: 7
; LENGTH: 669
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: human G-protein coupled receptor (GPCR) B4 amino
; US-10-096-144-7

Query Match 79.1%; Score 3514; DB 14; Length 669;
Best Local Similarity 98.4%; Pred. No. 0;
Matches 666; Conservative 1; Mismatches 2; Indels 8; Gaps 2;

Oy 162 ITTSAISDELDRDKRFPALLRTTPSADHVEAMVOLMHPFRMWIIVLVSSDTYGRDNGQ 221
Db 1 ITTSAISDELDRDKRFPALLRTTPSADHVEAMVOLMHPFRMWIIVLVSSDTYGRDNGQ 60
Oy 222 LIGERVARDDICIAFOETLPTLOPNOMTSEERORLVTIVDKLOOSTARVVVFPDLTL 281
Db 61 LIGERVARDDICIAFOETLPTLOPNOMTSEERORLVTIVDKLOOSTARVVVFPDLTL 120
Oy 282 YHFEVEVLKONFTGAVMIASBSMAIDPVLNHLTELGLGTFLGTTIOSVPIPGSEPREW 341
Db 121 YHFEVEVLKONFTGAVMIASBSMAIDPVLNHLTELGLGTFLGTTIOSVPIPGSEPREW 180
Oy 342 GPQAGPPLSRTSOSYTCNOECNCLNATLSFNTILRLSGERVVYSYSAVVAHALHS 401
Db 181 GPQAGPPLSRTSOSYTCNOECNCLNATLSFNTILRLSGERVVYSYSAVVAHALHS 240
Oy 402 LIGCDKSTCTKRVVPMQLEELIKVNFLLDHOIFPDPOGDVALHLEIVOMQDRSQNP 461
Db 241 LIGCDKSTCTKRVVPMQLEELIKVNFLLDHOIFPDPOGDVALHLEIVOMQDRSQNP 300
Oy 462 FOSVASYPLQROLKNIODISHTVNTIIPMSCKRQSGQKKKPVGIVHCCFECIDCL 521
Db 301 FOSVASYPLQROLKNIK-TSLHTVNTIIPMSCKRQSGQKKKPVGIVHCCFECIDCL 359
Oy 522 PGTFINTEDEYECACPNNEWSYOSETSCFKROLVLEHMEAPTIIVALLAALGFLSTL 581
Db 360 PGTFINTEDEYECACPNNEWSYOSETSCFKROLVLEHMEAPTIIVALLAALGFLSTL 412
Oy 582 AILVIFMRHFOPTIVRSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCROALFPLCF 641
Db 413 AILVIFMRHFOPTIVRSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCROALFPLCF 472

Oy 642 TICISCIASVFOIVCAFKMASREPRAYSYWVRQGYVSMAPITVLKMWIIVIGMLARP 701
Db 473 TICISCIASVFOIVCAFKMASREPRAYSYWVRQGYVSMAPITVLKMWIIVIGMLARP 532
Oy 702 QSHRTPDDPKITIVSCNPNYRNSLLFNSTLDSLVGFSFAYMKELEPTNNEAKFI 761
Db 533 QSHRTPDDPKITIVSCNPNYRNSLLFNSTLDSLVGFSFAYMKELEPTNNEAKFI 592
Oy 762 TLSMTFFYTSVSLCTFMSASVGLVTVTLVNLALISLQYFGPKCYMILFYPERN 821
Db 593 TLSMTFFYTSVSLCTFMSASVGLVTVTLVNLALISLQYFGPKCYMILFYPERN 652
Oy 822 TPAYNSMIOGYTMRD 838
Db 653 TPAYNSMIOGYTMRD 669

RESULT 9

US-10-225-567A-683
; Sequence 683, Application US/10225567A
; Publication No. US20030113798A1
; GENERAL INFORMATION:
; APPLICANT: Lifespan Biociences
; APPLICANT: Brown, Joseph P.
; APPLICANT: Burmer, Glenn C.
; APPLICANT: Roush, Christine L.
; TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
; FILE REFERENCE: 1920-4-4
; CURRENT APPLICATION NUMBER: US/10/225,567A
; PRIOR FILING DATE: 2001-12-19
; PRIOR APPLICATION NUMBER: 60/257,144
; PRIOR FILING DATE: 2000-12-19
; NUMBER OF SEQ ID NOS: 2292
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 683
; LENGTH: 669
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-10-225-567A-683

Query Match 79.1%; Score 3514; DB 14; Length 669;
Best Local Similarity 98.4%; Pred. No. 0;
Matches 666; Conservative 1; Mismatches 2; Indels 8; Gaps 2;

Oy 162 ITTSAISDELDRDKRFPALLRTTPSADHVEAMVOLMHPFRMWIIVLVSSDTYGRDNGQ 221
Db 1 ITTSAISDELDRDKRFPALLRTTPSADHVEAMVOLMHPFRMWIIVLVSSDTYGRDNGQ 60
Oy 222 LIGERVARDDICIAFOETLPTLOPNOMTSEERORLVTIVDKLOOSTARVVVFPDLTL 281
Db 61 LIGERVARDDICIAFOETLPTLOPNOMTSEERORLVTIVDKLOOSTARVVVFPDLTL 120
Oy 282 YHFEVEVLKONFTGAVMIASBSMAIDPVLNHLTELGLGTFLGTTIOSVPIPGSEPREW 341
Db 121 YHFEVEVLKONFTGAVMIASBSMAIDPVLNHLTELGLGTFLGTTIOSVPIPGSEPREW 180
Oy 342 GPQAGPPLSRTSOSYTCNOECNCLNATLSFNTILRLSGERVVYSYSAVVAHALHS 401
Db 181 GPQAGPPLSRTSOSYTCNOECNCLNATLSFNTILRLSGERVVYSYSAVVAHALHS 240
Oy 402 LIGCDKSTCTKRVVPMQLEELIKVNFLLDHOIFPDPOGDVALHLEIVOMQDRSQNP 461
Db 241 LIGCDKSTCTKRVVPMQLEELIKVNFLLDHOIFPDPOGDVALHLEIVOMQDRSQNP 300
Oy 462 FOSVASYPLQROLKNIODISHTVNTIIPMSCKRQSGQKKKPVGIVHCCFECIDCL 521
Db 301 FOSVASYPLQROLKNIK-TSLHTVNTIIPMSCKRQSGQKKKPVGIVHCCFECIDCL 359
Oy 522 PGTFINTEDEYECACPNNEWSYOSETSCFKROLVLEHMEAPTIIVALLAALGFLSTL 581
Db 360 PGTFINTEDEYECACPNNEWSYOSETSCFKROLVLEHMEAPTIIVALLAALGFLSTL 412
Oy 582 AILVIFMRHFOPTIVRSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCROALFPLCF 641

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Db      413  |||||ALIVFMHFRFTPIRSAGPMCFMLTLTLVAWVVPYVGPBKSTCLGQALFPLCF 472
Qy      642  |||||TICICIAVRSFOIYCAFMASRFPRAVSYWRYGQPIYSMAFITLKVIVIGMLARP 701
Db      473  |||||TICICIAVRSFOIYCAFMASRFPRAVSYWRYGQPIYSMAFITLKVIVIGMLARP 532
Qy      702  |||||OSHPRTDPPDKITTVSCNPNRNSLLEFNTSLDLLSVGFSFAVWKGELPTNNEAKFI 761
Db      533  |||||OSHPRTDPPDKITTVSCNPNRNSLLEFNTSLDLLSVGFSFAVWKGELPTNNEAKFI 592
Qy      762  |||||TLSMTFYFTSSVSLCTFMSAGSVLVTIVDLVTVLNLALSLGFGPKCYMILFPERN 821
Db      593  |||||TLSMTFYFTSSVSLCTFMSAGSVLVTIVDLVTVLNLALSLGFGPKCYMILFPERN 652
Qy      822  |||||TPAYNSMTQGYTMRD 838
Db      653  |||||TPAYNSMTQGYTMRD 669

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RESULT 10

```

US-09-927-315-7
; Sequence 7, Application US/09927315
; Publication No. US20030040045A1
; GENERAL INFORMATION:
; APPLICANT: Ryba, Charles S.
; APPLICANT: Nelson, Greg
; APPLICANT: Hoon, Mark A.
; APPLICANT: Chandrasekar, Jayaram
; APPLICANT: Zhang, Yifeng
; APPLICANT: The Regents of the University of California
; APPLICANT: The Government of the United States of America
; APPLICANT: as represented by the Secretary of the
; TITLE OF INVENTION: Mammalian Sweet Taste Receptors
; FILE REFERENCE: 02307E-120110US
; CURRENT APPLICATION NUMBER: US/09/927,315
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: US 60/302,898
; PRIOR FILING DATE: 2001-07-03
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 7
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURE:
; OTHER INFORMATION: rat T1R2 sweet taste receptor
US-09-927-315-7

```

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Query Match      72.1%; Score 3203.5; DB 10; Length 843;
Best Local Similarity 70.6%; Pred. No. 1.1e-293;
Matches 595; Conservative 108; Mismatches 133; Indels 7; Gaps 4;

Qy      1  |||||MEPRAKTICSLFLLMLVALEP---AENSDFYLPDGLLGLSLAHNMGIYHNLQVP 57
Db      1  |||||MEPRARTICLLSLHLVLPKPGKLYENSDFHLAGDVLGLGFTLHANVKSISHLSTLYVP 60
Qy      58  |||||MCKEVEKVIYGNLQAMRFAVEEINNDGSLPGVLLGYEIVDVCYSNNQPVLYFLAH 117
Db      61  |||||KNEFTMKVLGYNLQAMRFAVEEINNDGSLPGVLLGYEIVDVCYSNNQPVLYFLAH 120
Qy      118  |||||EDNLLPIQEDSYNYSRVAIVGPDNSSEVMTVAFLSLPQLITYAISDELDRKVF 177
Db      121  |||||DDDLPIILKDYQYMPHVAVVIGPDNSSESAITVSNLSHFLIPQITYAISDKLDRKF 180
Qy      178  |||||PALLRTPSADHVMAMVQMLHFRNMWITIVSSPTVGRDGLGGEVAR-RDICTAF 236
Db      181  |||||PSMLKRVPSATHIEMVQMLHFRNMWITIVSSPTVGRDGLGGEVAR-RDICTAF 240
Qy      237  |||||QETLPTLQPNQWMTSEERQRLVTIVDKLQOSTARVVVVSPDLTYHFNEVLRQFTGA 296

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Db      241  |||||QEVLPPIESSQVMSSEBQROLNIDLKLRTSARVVVSEBELSYSEFHEVLRNNTGF 300
Qy      297  |||||VWASESALIDPVLANHTELGLGFTIGITQSVPIPESEFEREMGPAGPPLSRTSQS 356
Db      301  |||||VWASESALIDPVLANHTELGLGFTIGITQSVPIPESEFEREMGPAGPPLSR 360
Qy      357  |||||YTCNQCNCNANLSEFNTIILRSGERVVSYSAVVAVNALSHLGLCDSSTCKRVVY 416
Db      361  |||||TTCNDQCAQNTTSSFNIIILISGERVVSYSAVVAVNALSHLGLCDSSTCKRVVY 420
Qy      417  |||||PMOLLEIWKVNFLLDHOIFPDPOGVALHLEIVQWQDRSONDFQSVASYPLQROUK 476
Db      421  |||||PMOLLEIWKVNFLLDHOIFPDPOGVALHLEIVQWQDRSONDFQSVASYPLQROUK 480
Qy      477  |||||NIQDISMTVNTLIPMSKSCRCQGGKKKRGVGHVCCFECIDCLPGFTLHTEDEYEQ 536
Db      481  |||||YINNVSWTYPNNVTVSVMSCKSCQGGKKKRGVGHVCCFECIDCLPGFTLHTEDEYEQ 540
Qy      537  |||||ACPNNEMSGOSETSCFKQLVLEWHEAPITIAVALALAGFLSTLAIIVIFMRHQPPIV 596
Db      541  |||||SCPGSMYSKNDITCFQRPFTLEWHEAPITIAVALALAGFLSTLAIIVIFMRHQPPIV 600
Qy      597  |||||RSAGGPMCFMLTLTLVAWVVPYVGPBKSTCLGQALFPLCTICICIAVRSFOIV 656
Db      601  |||||RSAGGPMCFMLTLTLVAWVVPYVGPBKSTCLGQALFPLCTICICIAVRSFOIV 660
Qy      657  |||||CAPKVASRFPRAVSYWRYGQPIYSMAFITLKVIVIGMLARQSHP--RTDPPDKI 714
Db      661  |||||CVFKVARLPASVYSMMVHGPVFAVITIAKVALVGNMLA-TTIPIGTDDDDPI 719
Qy      715  |||||TIVSCPNVRSNLFTSLDLLSVGFSFAVWKGELPTNNEAKFITLSMTFYFTSSVS 774
Db      720  |||||MILSCHPRYRGLEFNTSMDDLSTVGFSAVWKGELPTNNEAKFITLSMTFYFTSSVS 779
Qy      775  |||||LCTFMSAGSVLVTIVDLVTVLNLALSLGFGPKCYMILFPERNTPAYNSMTQGYT 834
Db      780  |||||LCTFMSAGSVLVTIVDLVTVLNLALSLGFGPKCYMILFPERNTPAYNSMTQGYT 839
Qy      835  |||||MR 837
Db      840  |||||MR 842

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RESULT 11

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US-10-124-598-1
; Sequence 1, Application US/10124598
; Publication No. US20020119526A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliott
; APPLICANT: Lindemeier, Uerger
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; FILE REFERENCE: 02307E-088720US
; CURRENT APPLICATION NUMBER: US/10/124,598
; CURRENT FILING DATE: 2002-04-16
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/361,631
; PRIOR FILING DATE: EARLIER FILING DATE: 1999-07-27
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/112,747
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURE:
; OTHER INFORMATION: rat G-protein coupled receptor (GPCR) B4 amino
; OTHER INFORMATION: acid sequence
US-10-124-598-1

```

```

Query Match      72.1%; Score 3203.5; DB 13; Length 843;

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Best Local Similarity 70.6%; Pred. No. 1.le-293;
Matches 595; Conservative 108; Mismatches 133; Indels 7; Gaps 4;

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QY 1 MGPRAKTICSLFFLLMVLAE---AENSDFYLPQDYLLGGLFSLHANKGIVHLNPLQVP 57
DB 1 MOPARTICLSLLHLVLPKPGKLVENSDFHLAGDYLGLFTLHANKVKSISHLSYLOVP 60
QY 58 MCKEYEVKIVGYNLQMAMRAVEBEINNDSSLLPGVLLGEIYDVYCSNNVOPVLYPLAH 117
DB 61 KNEFTMKVLYGYNLQMAMRAVEBEINNDSSLLPGVLLGEIYDVYCSNNVOPVLYPLAH 120
QY 118 EDNLPIQEDYNSYISRVAVVIGPNSSESVMTVANFLSLFTLPQITYSALSDLRDQVRF 177
DB 121 DDDLPLIKDYSQYMPHVAVVIGPNSSESVMTVANFLSLFTLPQITYSALSDLRDQVRF 180
QY 178 PALRTTPSADHHEAVVQLMHLFRNMNIIVLSSDPTVGRDQGLGERVAR-RDICIAP 236
DB 181 PSMRTVPSATHHIEAVVQLMHLFRNMNIIVLSSDPTVGRDQGLGERVAR-RDICIAP 240
QY 237 QETLFTLPQNMNTSEERQRLVTIVDKLQOSTARVVVSPDLTLVHFENEVLRONFTGA 296
DB 241 QEVLPPIESSQVMSRSEERQRLVDLKLRTSARVVVSPDLTLVHFENEVLRONFTGA 300
QY 297 VMAISESWAIDPVHLNLTGHTGFTLGTITQSPFIPGSEFERMGPOAPPPLSRISQS 356
DB 301 VMAISESWAIDPVHLNLTGHTGFTLGTITQSPFIPGSEFERMGPOAPPPLSRISQS 360
QY 357 YTCNOECNCLNATLSFNTILRLSEGERVYVSAYVAVAHALSHLSDGCKSTCKRVV 416
DB 361 TTCNOECNCLNATLSFNTILRLSEGERVYVSAYVAVAHALSHLSDGCKSTCKRVV 420
QY 417 PMOLLEIWKVNFLLDHOIFPDQGVALLHLEIVQWQMSQNPQSVASVYPLQRLK 476
DB 421 PMOLLEIWKVNFLLDHOIFPDQGVALLHLEIVQWQMSQNPQSVASVYPLQRLK 480
QY 477 NIQDISHTVNTIIPMSCKRQSGQKKGIVHVCCEECIDCLPGLFNLHTEBEYEQ 536
DB 481 YINNVSWYTPNNVTPVMSCKSKQOPGQMKSVGLHPCCEFCIDCMPGTYLNRSADEFNCL 540
QY 537 ACPNNEWSYQSETSCEFRLQVLEMEHAPTIYVALLAALGFLSTLAIIVIFMRHPQPIV 596
DB 541 SCPSGMSYKNDITCFQRRPFTLEMEHAPTIYVALLAALGFLSTLAIIVIFMRHPQPIV 600
QY 597 RSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCQALPPLCTICISCIARVSQIV 656
DB 601 RSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCQALPPLCTICISCIARVSQIV 660
QY 657 CAFKMASRFPRAYSYWRVYQGPVYSMAFIVLAKMIVVYIGMLARQSHR--RTDPDPKI 714
DB 661 CAFKMASRFPRAYSYWRVYQGPVYSMAFIVLAKMIVVYIGMLARQSHR--RTDPDPKI 719
QY 715 TIVSCNPNYRNSILFNTSLDLSLVGFSFAYMGKELPTNVEAKFITLSTMTFTYSVS 774
DB 720 MILSHPNYRNGLPNTSMDLISLVGFSFAYMGKELPTNVEAKFITLSTMTFTYSVS 779
QY 775 LCTFMSAVSGVLYTIVDLVTVLNLALISLGYFGKCYMLFYPEBRNTPAYFNSMIQYT 834
DB 780 LCTFMSAVSGVLYTIVDLVTVLNLALISLGYFGKCYMLFYPEBRNTPAYFNSMIQYT 839
QY 835 MRR 837
DB 840 MRR 842
```

RESULT 12
US-10-096-144-1

; Sequence 1, Application US/10096144
; Publication No. US20030022288A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Adler, Jon Elliot
; APPLICANT: Lindemeier, Juergen
; APPLICANT: The Regents of the University of California

; TITLE OF INVENTION: Nucleic Acids Encoding a G-Protein Coupled Receptor
; FILE OF INVENTION: Involved in Sensory Transduction
; FILE REFERENCE: 02307E-088720US
; CURRENT APPLICATION NUMBER: US/10/096.144
; PRIOR FILING DATE: 2002-03-08
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/361.631
; PRIOR FILING DATE: EARLIER FILING DATE: 1999-07-27
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/095.464
; PRIOR FILING DATE: EARLIER FILING DATE: 1998-07-28
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US 60/112.747
; PRIOR FILING DATE: EARLIER FILING DATE: 1998-12-17
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 1
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
; FEATURE:
; OTHER INFORMATION: rat G-protein coupled receptor (GPCR) B4 amino
; OTHER INFORMATION: acid sequence
US-10-096-144-1

Query Match 72.1%; Score 3203.5; DB 14; Length 843;

Best Local Similarity 70.6%; Pred. No. 1.le-293;
Matches 595; Conservative 108; Mismatches 133; Indels 7; Gaps 4;

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QY 1 MGPRAKTICSLFFLLMVLAE---AENSDFYLPQDYLLGGLFSLHANKGIVHLNPLQVP 57
DB 1 MOPARTICLSLLHLVLPKPGKLVENSDFHLAGDYLGLFTLHANKVKSISHLSYLOVP 60
QY 58 MCKEYEVKIVGYNLQMAMRAVEBEINNDSSLLPGVLLGEIYDVYCSNNVOPVLYPLAH 117
DB 61 KNEFTMKVLYGYNLQMAMRAVEBEINNDSSLLPGVLLGEIYDVYCSNNVOPVLYPLAH 120
QY 118 EDNLPIQEDYNSYISRVAVVIGPNSSESVMTVANFLSLFTLPQITYSALSDLRDQVRF 177
DB 121 DDDLPLIKDYSQYMPHVAVVIGPNSSESVMTVANFLSLFTLPQITYSALSDLRDQVRF 180
QY 178 PALRTTPSADHHEAVVQLMHLFRNMNIIVLSSDPTVGRDQGLGERVAR-RDICIAP 236
DB 181 PSMRTVPSATHHIEAVVQLMHLFRNMNIIVLSSDPTVGRDQGLGERVAR-RDICIAP 240
QY 237 QETLFTLPQNMNTSEERQRLVTIVDKLQOSTARVVVSPDLTLVHFENEVLRONFTGA 296
DB 241 QEVLPPIESSQVMSRSEERQRLVDLKLRTSARVVVSPDLTLVHFENEVLRONFTGA 300
QY 297 VMAISESWAIDPVHLNLTGHTGFTLGTITQSPFIPGSEFERMGPOAPPPLSRISQS 356
DB 301 VMAISESWAIDPVHLNLTGHTGFTLGTITQSPFIPGSEFERMGPOAPPPLSRISQS 360
QY 357 YTCNOECNCLNATLSFNTILRLSEGERVYVSAYVAVAHALSHLSDGCKSTCKRVV 416
DB 361 TTCNOECNCLNATLSFNTILRLSEGERVYVSAYVAVAHALSHLSDGCKSTCKRVV 420
QY 417 PMOLLEIWKVNFLLDHOIFPDQGVALLHLEIVQWQMSQNPQSVASVYPLQRLK 476
DB 421 PMOLLEIWKVNFLLDHOIFPDQGVALLHLEIVQWQMSQNPQSVASVYPLQRLK 480
QY 477 NIQDISHTVNTIIPMSCKRQSGQKKGIVHVCCEECIDCLPGLFNLHTEBEYEQ 536
DB 481 YINNVSWYTPNNVTPVMSCKSKQOPGQMKSVGLHPCCEFCIDCMPGTYLNRSADEFNCL 540
QY 537 ACPNNEWSYQSETSCEFRLQVLEMEHAPTIYVALLAALGFLSTLAIIVIFMRHPQPIV 596
DB 541 SCPSGMSYKNDITCFQRRPFTLEMEHAPTIYVALLAALGFLSTLAIIVIFMRHPQPIV 600
QY 597 RSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCQALPPLCTICISCIARVSQIV 656
DB 601 RSAGGPMCFMLTLLVAVVVPVYVGPVKVSTCLCQALPPLCTICISCIARVSQIV 660
QY 657 CAFKMASRFPRAYSYWRVYQGPVYSMAFIVLAKMIVVYIGMLARQSHR--RTDPDPKI 714
DB 661 CAFKMASRFPRAYSYWRVYQGPVYSMAFIVLAKMIVVYIGMLARQSHR--RTDPDPKI 719
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QY 58 MCKEYEVKIVGYNLMQAMRFAVEEINNDSLLPGVLLGEYIVDYCYISNNVOPVLYFLAH 117
DB 61 KCNEBTMVLGVGNLMQAMRFAVEEINNDSLLPGVLLGEYIVDYCYISNNVOPVLYFLAH 120
QY 118 EDNLLPIQEDBSYNTISRVAIVIGPNSSESVMTVANFLSLFLPQITYSASIDELDKYRF 177
DB 121 DDDLLPIKDYSOQYMPHVAVIGPNSSESAIVSNILSHFLIPQITYSASIDELDKYRF 180
QY 178 PALARTPSADHVEAMVQMLMFRMNNITIVVSDTYGRDNGOLLSGRVAR-RDICIATF 236
DB 181 PSMLTRVTSATHTIEMAVQMLMFRMNNITIVVSDTYGRDNGOLLSGRVAR-RDICIATF 240
QY 237 QETLPTLOPNQNMNTSEERQRLVTVDKLQOSTARVVVSPDLTYHFENEVLARQNTGA 296
DB 241 QEVLPIRESSQVMRSEBQRLDNILDKLRTSARVVVSPDLTYHFENEVLARQNTGA 300
QY 297 VWIASESAIDPVLANLTBLGHLGTLGTLTTSQVPIPGSEPREMGPOAGPPPLSRTSQS 356
DB 301 VWIASESAIDPVLANLTBLGHLGTLGTLTTSQVPIPGSEPREMGPOAGPPPLSRTSQS 360
QY 357 YTCNOECNCLNATLSFNTILRLSGERVVYSVYSAVVAHALHSLGCDSTCKRXYV 416
DB 361 TTCNQDCACLNATLSFNTILRLSGERVVYSVYSAVVAHALHSLGCDSTCKRXYV 420
QY 417 PMQLLEIWKVNFLLDHOIFPDQGVVALHLEIVQOMQDRSQNPQSVASYPLQRLK 476
DB 421 PMQLLEIWKVNFLLDHOIFPDQGVVALHLEIVQOMQDRSQNPQSVASYPLQRLK 480
QY 477 NIODISWHTVNTTIPMSGSKRCQSGQKKPVGIVHVCCEPICDCLPGFLNHTEDEXEQ 536
DB 481 YINNVSWYTPNNVTPVSMCSKSCOPGOMKSVGLHPCCEPICDCLPGFLNHTEDEXEQ 540
QY 537 ACPNNEWSYQSESCFRQOLVFLMEHBAPTIAVALAALGFLSTALIVIFMRHFOPTIV 596
DB 541 SCPSMWSYKNDITCFQRPTFLEMEHBAPTIAVALAALGFLSTALIVIFMRHFOPTIV 600
QY 597 RSAGGPMCFMLTLVLAVMVVPYVGPVKYSTCLCQALPPLCFTICISCIARVFOIV 656
DB 601 RSAGGPMCFMLTLVLAVMVVPYVGPVKYSTCLCQALPPLCFTICISCIARVFOIV 660
QY 657 CAFKASFPFPAVYSWVYQGPYSMATIVLAKVIYVIGMLARQSPH--RTDDEPKI 714
DB 661 CVFKARLPSPASVFMKRYGPPVAFITAIKVALVVGMMLA--TTINPIGITDDEPKI 719
QY 715 TIVSCNPNRSLFNTSLDILLVSGFSPAYMGEKLPNTNBEAKFILTSMTEYTSVS 774
DB 720 MILSCHPNRGNGLFNTSMDDLVLGFSFAYMGKELPTNNEAKFILTSMTEYTSVS 779
QY 775 LCTFMSAYSGVLTVLVDLVTVLNLALASLGFGPKCYMILFPYPERNTPAYFNSMIOGYT 834
DB 780 LCTFMSAYSGVLTVLVDLVTVLNLALASLGFGPKCYMILFPYPERNTPAYFNSMIOGYT 839
QY 835 MRR 837
DB 840 MRR 842

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RESULT 15
US-10-179-373-17
; Sequence 17, Application US/10179373
; Publication No. US20030232407A1
; GENERAL INFORMATION:
; APPLICANT: ZOLLER, MARK
; APPLICANT: LI, XINADONG
; APPLICANT: STASZEMSKI, LENA
; APPLICANT: O'CONNELL, SHAWN
; APPLICANT: ZOZULYA, SERGEY
; APPLICANT: ADLER, JON
; APPLICANT: XU, HONG
; APPLICANT: ECHEVERRI, FERNANDO
; TITLE OF INVENTION: TIR HETERO-OLIGOMERIC TASTE RECEPTORS AND CELL LINES
; THAT EXPRESS SAID RECEPTORS AND USE THEREOF

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; TITLE OF INVENTION: IDENTIFICATION OF TASTE COMPOUNDS
; FILE REFERENCE: 078003-0291566
; CURRENT APPLICATION NUMBER: US/10/179, 373
; CURRENT FILING DATE: 2002-06-26
; PRIOR APPLICATION NUMBER: 60/300, 434
; PRIOR FILING DATE: 2001-06-26
; PRIOR APPLICATION NUMBER: 60/304, 749
; PRIOR FILING DATE: 2001-07-13
; PRIOR APPLICATION NUMBER: 60/310, 493
; PRIOR FILING DATE: 2001-08-08
; PRIOR APPLICATION NUMBER: 60/331, 771
; PRIOR FILING DATE: 2001-11-21
; PRIOR APPLICATION NUMBER: 60/339, 472
; PRIOR FILING DATE: 2001-12-14
; PRIOR APPLICATION NUMBER: 60/372, 090
; PRIOR FILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: 60/374, 143
; NUMBER OF SEQ ID NOS: 19
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 17
; LENGTH: 843
; TYPE: PRT
; ORGANISM: Rattus sp.
US-10-179-373-17

Query Match 72.1%; Score 3203.5; DB 15; Length 843;
Best Local Similarity 70.6%; Pred. No. 1.1e-293;
Matches 595; Conservative 108; Mismatches 133; Indels 7; Gaps 4;

QY 1 MGPRAKTCISLFLMLWYLAEB--AENSDFYLPEDYLLGLGFLSHANKGIVHLNPLQVP 57
DB 1 MGPOARTCLISLHLVLPKRGKLVENSDFHLADYLLGLGFLSHANKGIVHLNPLQVP 60
QY 58 MCKEYEVKIVGYNLMQAMRFAVEEINNDSLLPGVLLGEYIVDYCYISNNVOPVLYFLAH 117
DB 61 KCNEBTMVLGVGNLMQAMRFAVEEINNDSLLPGVLLGEYIVDYCYISNNVOPVLYFLAH 120
QY 118 EDNLLPIQEDBSYNTISRVAIVIGPNSSESVMTVANFLSLFLPQITYSASIDELDKYRF 177
DB 121 DDDLLPIKDYSOQYMPHVAVIGPNSSESAIVSNILSHFLIPQITYSASIDELDKYRF 180
QY 178 PALARTPSADHVEAMVQMLMFRMNNITIVVSDTYGRDNGOLLSGRVAR-RDICIATF 236
DB 181 PSMLTRVTSATHTIEMAVQMLMFRMNNITIVVSDTYGRDNGOLLSGRVAR-RDICIATF 240
QY 237 QETLPTLOPNQNMNTSEERQRLVTVDKLQOSTARVVVSPDLTYHFENEVLARQNTGA 296
DB 241 QEVLPIRESSQVMRSEBQRLDNILDKLRTSARVVVSPDLTYHFENEVLARQNTGA 300
QY 297 VWIASESAIDPVLANLTBLGHLGTLGTLTTSQVPIPGSEPREMGPOAGPPPLSRTSQS 356
DB 301 VWIASESAIDPVLANLTBLGHLGTLGTLTTSQVPIPGSEPREMGPOAGPPPLSRTSQS 360
QY 357 YTCNOECNCLNATLSFNTILRLSGERVVYSVYSAVVAHALHSLGCDSTCKRXYV 416
DB 361 TTCNQDCACLNATLSFNTILRLSGERVVYSVYSAVVAHALHSLGCDSTCKRXYV 420
QY 417 PMQLLEIWKVNFLLDHOIFPDQGVVALHLEIVQOMQDRSQNPQSVASYPLQRLK 476
DB 421 PMQLLEIWKVNFLLDHOIFPDQGVVALHLEIVQOMQDRSQNPQSVASYPLQRLK 480
QY 477 NIODISWHTVNTTIPMSGSKRCQSGQKKPVGIVHVCCEPICDCLPGFLNHTEDEXEQ 536
DB 481 YINNVSWYTPNNVTPVSMCSKSCOPGOMKSVGLHPCCEPICDCLPGFLNHTEDEXEQ 540
QY 537 ACPNNEWSYQSESCFRQOLVFLMEHBAPTIAVALAALGFLSTALIVIFMRHFOPTIV 596
DB 541 SCPSMWSYKNDITCFQRPTFLEMEHBAPTIAVALAALGFLSTALIVIFMRHFOPTIV 600
QY 597 RSAGGPMCFMLTLVLAVMVVPYVGPVKYSTCLCQALPPLCFTICISCIARVFOIV 656
DB 601 RSAGGPMCFMLTLVLAVMVVPYVGPVKYSTCLCQALPPLCFTICISCIARVFOIV 660

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us-09-927-315-15.rapb

Page 1

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run on: May 11, 2004, 15:27:42 ; Search time 41.336 Seconds
(without alignments)
5720.583 Million cell updates/sec

Title: US-09-927-315-15

Perfect score: 4524

Sequence: 1 MLGPVILGLSLMALHPGTG.....GPGDAQGNQDNTGNQKHE 852

Scoring table:

BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1140673 seqs, 27756755 residues

Total number of hits satisfying chosen parameters: 1140673

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database :

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. Published Applications AA:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	4524	100.0	852	10	US-09-927-315-15
3	4524	100.0	852	14	US-10-035-045-4
4	4524	100.0	852	14	US-10-190-417-15
5	4524	100.0	936	15	US-10-292-798-602
6	4517	99.8	852	12	US-10-188-186-148
7	4512	99.7	852	9	US-09-796-338A-14
8	4512	99.7	852	14	US-10-261-482-2
9	4512	99.7	852	14	US-10-283-837-14
10	4512	99.7	852	14	US-10-145-586-14
11	4512	99.7	852	15	US-10-407-079-90
12	4512	99.7	852	15	US-10-179-373-7
13	4509	99.7	852	16	US-10-280-163A-5
14	4500	99.5	850	10	US-09-799-629-4
15	4500	99.5	850	14	US-10-190-417-30

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17	4490	99.2	850	14	US-10-246-785-7	Sequence 7, Appli
18	4474	98.9	894	15	US-10-436-715-6	Sequence 6, Appli
19	4461	98.6	854	14	US-10-159-339-2	Sequence 2, Appli
20	4461	98.6	854	15	US-10-436-715-86	Sequence 86, Appli
21	4365.5	96.5	829	15	US-10-436-715-4	Sequence 4, Appli
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23	4002	88.5	771	14	US-10-225-567A-593	Sequence 593, App
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27	3255	71.9	858	14	US-10-035-045-14	Sequence 14, Appli
28	3255	71.9	858	14	US-10-246-785-9	Sequence 9, Appli
29	3255	71.9	858	14	US-10-190-417-25	Sequence 25, Appli
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33	3229	71.4	858	14	US-10-190-417-23	Sequence 23, Appli
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35	3227.5	71.3	858	10	US-09-927-315-18	Sequence 18, Appli
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37	3221.5	71.2	858	10	US-09-927-315-20	Sequence 20, Appli
38	3221.5	71.2	858	14	US-10-190-417-20	Sequence 20, Appli
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45	1208.5	26.7	840	14	US-10-190-417-1	Sequence 1, Appli

ALIGNMENTS

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RESULT 1
US-09-897-427A-6
; Sequence 6, Application US/09897427A
; Patent No. US2002016042A1
; GENERAL INFORMATION:
; APPLICANT: ADLER, JON ELLIOT
; APPLICANT: LI, XIADONG
; APPLICANT: STAZESKI, LENA
; APPLICANT: XU, HONG
; APPLICANT: EHEVERRI, FERNANDO
; TITLE OF INVENTION: T1R HETERO-OLIGOMERIC TASTE RECEPTORS
; FILE REFERENCE: 078003-0282558
; CURRENT APPLICATION NUMBER: US/09/897,427A
; CURRENT FILING DATE: 2001-07-03
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; TYPE: PRT
; LENGTH: 852
; ORGANISM: Homo sapiens
US-09-897-427A-6
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Query Match 100.0%; Score 4524; DB 9; Length 852;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 852; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 2

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US-09-927-315-15
; Sequence 15, Application US/09927315
; Publication No. US20030040045A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Ryba, Nicholas J. P.
; APPLICANT: Nelson, Greg
; APPLICANT: Hoon, Mark A.
; APPLICANT: Chandrasekar, Jayaram
; APPLICANT: Zhang, Yifeng
; APPLICANT: The Regents of the University of California
; APPLICANT: as represented by the United States of America
; APPLICANT: Department of Health and Human Services
; TITLE OF INVENTION: Mammalian Sweet Taste Receptors
; FILE REFERENCE: 02307E-1201I0US
; CURRENT APPLICATION NUMBER: US/09/927,315
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: US 60/302,898
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 15
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; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: human T1R3 sweet taste receptor
US-09-927-315-15
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Query Match 100.0%; Score 4524; DB 10; Length 852;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 852; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 3

Wed May 12 10:42:53 2004

us-09-927-315-15.rapb

Page 3

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; Sequence 4, Application US/10035045
; Publication No. US20030054448A1
; GENERAL INFORMATION:
; APPLICANT: ADLER, JON ELLIOT
; APPLICANT: LI, XIADONG
; APPLICANT: STASZEWSKI, LENA
; APPLICANT: O'CONNELL, SHAWN
; APPLICANT: ZOZULYA, SERGEY
; TITLE OF INVENTION: TIR TASTE RECEPTORS AND GENES ENCODING SAME
; FILE REFERENCE: 078003-0280681
; CURRENT APPLICATION NUMBER: US/10/035,045
; PRIOR FILING DATE: 2002-01-03
; PRIOR APPLICATION NUMBER: 60/259,227
; PRIOR FILING DATE: 2001-01-03
; PRIOR APPLICATION NUMBER: 60/284,547
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-035-045-4

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Beet Local Similarity 100.0%; Pred. No. 0;
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DB      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
QY      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
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DB      301  EAMLTSDLVMGILPGMAQKGTIVGLFQORGALHEFPQYVYKTHALATDPAFCALGERREG 360
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RESULT 4
US-10-190-417-15
; Sequence 15, Application US/10190417
; Publication No. US20030166137A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Ryba, Nicholas J. P.
; APPLICANT: Chandrasekar, Jayaram
; APPLICANT: Hoon, Mark A.
; APPLICANT: Nelson, Greg
; APPLICANT: Zhang, Yifeng
; APPLICANT: The Regents of the University of California
; APPLICANT: The Government of the United States of America
; APPLICANT: as represented by the Secretary of the
; APPLICANT: Department of Health and Human Services
; TITLE OF INVENTION: Mammalian Sweet and Amino Acid Heterodimeric Taste
; FILE REFERENCE: 023078-120130US
; CURRENT APPLICATION NUMBER: US/10/190,417
; PRIOR FILING DATE: 2002-11-14
; PRIOR APPLICATION NUMBER: US 60/302,898
; PRIOR FILING DATE: 2001-07-03
; PRIOR APPLICATION NUMBER: US 09/927,315
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: US 60/358,925
; PRIOR FILING DATE: 2002-02-22
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 15
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: human T1R3
US-10-190-417-15

Query Match      100.0%; Score 4524; DB 14; Length 852;
Beet Local Similarity 100.0%; Pred. No. 0;
Matches 852; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1  MGGPAVLGLSLWALLHPGTGAPLCLISQQLRMKGDVYLGLPFLGAEBAAGLRSTRPSSP 60
DB      1  MGGPAVLGLSLWALLHPGTGAPLCLISQQLRMKGDVYLGLPFLGAEBAAGLRSTRPSSP 60
QY      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
DB      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
QY      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
DB      61  VCTRSSNGCLMALMKAAVEEINKSDLLPGLRGYDLPTCCSEPPVAMKPSLMFLAKA 120
QY      121  GSRDIAACNTYQYOPRYLAVIGPHSSSELAVYTGKFFSLMPQVSYGASMEILLSARETF 180
DB      121  GSRDIAACNTYQYOPRYLAVIGPHSSSELAVYTGKFFSLMPQVSYGASMEILLSARETF 180
QY      121  GSRDIAACNTYQYOPRYLAVIGPHSSSELAVYTGKFFSLMPQVSYGASMEILLSARETF 180
DB      121  GSRDIAACNTYQYOPRYLAVIGPHSSSELAVYTGKFFSLMPQVSYGASMEILLSARETF 180
QY      181  PSFRTVSDRVQULTAAELLQEFGMNVAAALGSDDEYRGQGLSIFSAALAAARGICIAHE 240
DB      181  PSFRTVSDRVQULTAAELLQEFGMNVAAALGSDDEYRGQGLSIFSAALAAARGICIAHE 240
```

QY 241 GLVPLPRADSRIGKVQDVHLHQVNQSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
DB 241 GLVPLPRADSRIGKVQDVHLHQVNQSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
QY 301 EAMLTSDLVNGLPGMAQMGTVLGFLORGAQLHEFPQYKTHLALATDPAFCGALGEREQ 360
DB 301 EAMLTSDLVNGLPGMAQMGTVLGFLORGAQLHEFPQYKTHLALATDPAFCGALGEREQ 360
QY 361 LEEDVVGORCPQDCITLQNVASAGLNHHQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
DB 361 LEEDVVGORCPQDCITLQNVASAGLNHHQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
QY 421 VKPQQLLENNYNTLFHVGLPLRFDSSGNVMEYDLKLMWQGSVPRLDVGRFNGSLRT 480
DB 421 VKPQQLLENNYNTLFHVGLPLRFDSSGNVMEYDLKLMWQGSVPRLDVGRFNGSLRT 480
QY 481 ERLKIRMTSDNQKPVSRCSRCQCEGOVRVKGFSGCCYDCVCEAGSYRONPDDIACTF 540
DB 481 ERLKIRMTSDNQKPVSRCSRCQCEGOVRVKGFSGCCYDCVCEAGSYRONPDDIACTF 540
QY 541 CGODEMSPERSTRCFRRRSRFLAMGEPAYLLLLLLLSTALGLVLAALGFVHHRDSPLVQ 600
DB 541 CGODEMSPERSTRCFRRRSRFLAMGEPAYLLLLLLLSTALGLVLAALGFVHHRDSPLVQ 600
QY 601 ASGGPLACFGVLGVLCLSVLLFPQOPSPARCIAQOPLSHLPLTGCLSTLFLQAAE1FV 660
DB 601 ASGGPLACFGVLGVLCLSVLLFPQOPSPARCIAQOPLSHLPLTGCLSTLFLQAAE1FV 660
QY 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAPPEVVTDMHMLPTEALV 720
DB 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAPPEVVTDMHMLPTEALV 720
QY 721 HCRTRSWVSFGLAHATNATLAFCLGTFVLVRSQPCYNBARGLTFAMLAFFITWVSFVP 780
DB 721 HCRTRSWVSFGLAHATNATLAFCLGTFVLVRSQPCYNBARGLTFAMLAFFITWVSFVP 780
QY 781 LLANVQVVLPRPAVQMGALLLCVGIILAAFHLPRLCYLLMRQPLNTPPEFLGGGPGDAQO 840
DB 781 LLANVQVVLPRPAVQMGALLLCVGIILAAFHLPRLCYLLMRQPLNTPPEFLGGGPGDAQO 840
QY 841 NDGNTGNOGKHE 852
DB 841 NDGNTGNOGKHE 852

RESULT 5
US-10-292-798-602
; Sequence 602, Application US/10292798
; Publication No. US20030235833A1
; GENERAL INFORMATION:
; APPLICANT: SUMA, MAKIKO
; APPLICANT: ASAI, KIYOSHI
; APPLICANT: AKIYAMA, YUTAKA
; APPLICANT: ABURATANI, HIROYUKI
; TITLE OF INVENTION: GANDOSINE TRIPHOSPHATE-BINDING PROTEIN COUPLED RECEPTORS
; FILE REFERENCE: 084335/166
; CURRENT APPLICATION NUMBER: US/10/292,798
; PRIOR FILING DATE: 2002-11-13
; PRIOR APPLICATION NUMBER: 10/017,161
; PRIOR FILING DATE: 2001-12-18
; PRIOR APPLICATION NUMBER: JP 2001-246789
; PRIOR FILING DATE: 2001-06-18
; NUMBER OF SEQ ID NOS: 2070
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 602
; LENGTH: 936
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-292-798-602

Query Match 100.0%; Score 4524; DB 15; Length 936;
Beet Local Similarity 100.0%; Pred. No. 0;
Matches 852; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MLGPAVILGSLMALHPEGTGAPLCLSQQLRMKQDYVLGGFLPGAEAEAGLRSTRPSSP 60
DB 1 MLGPAVILGSLMALHPEGTGAPLCLSQQLRMKQDYVLGGFLPGAEAEAGLRSTRPSSP 60
QY 61 VCTRFSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFDTCSEPVVANKPSIMPLAKA 120
DB 61 VCTRFSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFDTCSEPVVANKPSIMPLAKA 120
QY 121 GSRDIAAYCNYTOVQPRVLAVIGPSSSELMYMGKFPSPFLMPQVSYGASMEILSARETF 180
DB 121 GSRDIAAYCNYTOVQPRVLAVIGPSSSELMYMGKFPSPFLMPQVSYGASMEILSARETF 180
QY 181 PSFPRTPSDRVOLTTAAAEILQEFGNMNVAAALGSDDEYGRQGLSIFGALAAAGICTAHE 240
DB 181 PSFPRTPSDRVOLTTAAAEILQEFGNMNVAAALGSDDEYGRQGLSIFGALAAAGICTAHE 240
QY 241 GLVPLPRADSRIGKVQDVHLHQVNQSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
DB 241 GLVPLPRADSRIGKVQDVHLHQVNQSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
QY 301 EAMLTSDLVNGLPGMAQMGTVLGFLORGAQLHEFPQYKTHLALATDPAFCGALGEREQ 360
DB 301 EAMLTSDLVNGLPGMAQMGTVLGFLORGAQLHEFPQYKTHLALATDPAFCGALGEREQ 360
QY 361 LEEDVVGORCPQDCITLQNVASAGLNHHQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
DB 361 LEEDVVGORCPQDCITLQNVASAGLNHHQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
QY 421 VKPQQLLENNYNTLFHVGLPLRFDSSGNVMEYDLKLMWQGSVPRLDVGRFNGSLRT 480
DB 421 VKPQQLLENNYNTLFHVGLPLRFDSSGNVMEYDLKLMWQGSVPRLDVGRFNGSLRT 480
QY 481 ERLKIRMTSDNQKPVSRCSRCQCEGOVRVKGFSGCCYDCVCEAGSYRONPDDIACTF 540
DB 481 ERLKIRMTSDNQKPVSRCSRCQCEGOVRVKGFSGCCYDCVCEAGSYRONPDDIACTF 540
QY 541 CGODEMSPERSTRCFRRRSRFLAMGEPAYLLLLLLLSTALGLVLAALGFVHHRDSPLVQ 600
DB 541 CGODEMSPERSTRCFRRRSRFLAMGEPAYLLLLLLLSTALGLVLAALGFVHHRDSPLVQ 600
QY 601 ASGGPLACFGVLGVLCLSVLLFPQOPSPARCIAQOPLSHLPLTGCLSTLFLQAAE1FV 660
DB 601 ASGGPLACFGVLGVLCLSVLLFPQOPSPARCIAQOPLSHLPLTGCLSTLFLQAAE1FV 660
QY 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAPPEVVTDMHMLPTEALV 720
DB 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAPPEVVTDMHMLPTEALV 720
QY 721 HCRTRSWVSFGLAHATNATLAFCLGTFVLVRSQPCYNBARGLTFAMLAFFITWVSFVP 780
DB 721 HCRTRSWVSFGLAHATNATLAFCLGTFVLVRSQPCYNBARGLTFAMLAFFITWVSFVP 780
QY 781 LLANVQVVLPRPAVQMGALLLCVGIILAAFHLPRLCYLLMRQPLNTPPEFLGGGPGDAQO 840
DB 781 LLANVQVVLPRPAVQMGALLLCVGIILAAFHLPRLCYLLMRQPLNTPPEFLGGGPGDAQO 840
QY 841 NDGNTGNOGKHE 852
DB 841 NDGNTGNOGKHE 852

RESULT 6
US-10-188-186-148
; Sequence 148, Application US/10188186
; Publication No. US20040029789A1
; GENERAL INFORMATION:
; APPLICANT: Anderson et al.
; TITLE OF INVENTION: NOVEL PROTEINS AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 21402-397C
; CURRENT APPLICATION NUMBER: US/10/188,186
; PRIOR FILING DATE: 2002-07-02
; PRIOR APPLICATION NUMBER: 60/303046

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; PRIOR FILING DATE: 2001-07-05
; PRIOR APPLICATION NUMBER: 60/360814
; PRIOR FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: 60/303828
; PRIOR FILING DATE: 2001-09-07
; PRIOR APPLICATION NUMBER: 60/323380
; PRIOR FILING DATE: 2001-09-19
; PRIOR APPLICATION NUMBER: 60/361133
; PRIOR FILING DATE: 2002-03-01
; PRIOR APPLICATION NUMBER: 60/304016
; PRIOR FILING DATE: 2001-07-09
; PRIOR APPLICATION NUMBER: 60/304502
; PRIOR FILING DATE: 2001-07-11
; PRIOR APPLICATION NUMBER: 60/305262
; PRIOR FILING DATE: 2001-07-13
; PRIOR APPLICATION NUMBER: 60/373881
; PRIOR FILING DATE: 2002-04-19
; PRIOR APPLICATION NUMBER: 60/305673
; PRIOR FILING DATE: 2001-07-16
; Remaining Prior Application data removed - See file Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 368
; SOFTWARE: Custom
; SEQ ID NO: 148
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-188-186-148

Query Match      99.8%; Score 4517; DB 12; Length 852;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1  MCGPAVLGLSLMALHPGTGAPLCLISQQLRMKGDIYVGLFPLGAEAEAGLRSRTRPSSP 60
DB      1  MCGPAVLGLSLMALHPGTGAPLCLISQQLRMKGDIYVGLFPLGAEAEAGLRSRTRPSSP 60
QY      61  VCTRSSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFPTCSEPPVAMKPSLMFLAKA 120
DB      61  VCTRSSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFPTCSEPPVAMKPSLMFLAKA 120
QY      121  GSRDIAAYCNTYOQPRVLAVIGPHSSBLAVTGFKEFSPFLMPQVSYGASMEILSARETF 180
DB      121  GSRDIAAYCNTYOQPRVLAVIGPHSSBLAVTGFKEFSPFLMPQVSYGASMEILSARETF 180
QY      181  PSFFPTVSDRYQLTAAAEELLQEFGMNVVAALGSDDEYGRQGLSTFSALAAARGICIAHE 240
DB      181  PSFFPTVSDRYQLTAAAEELLQEFGMNVVAALGSDDEYGRQGLSTFSALAAARGICIAHE 240
QY      241  GLVPLPRADDSRLGKVQDVLLHOVNOSSVQVLLFPASVHAHAALFNYSISSRLSPRYWAS 300
DB      241  GLVPLPRADDSRLGKVQDVLLHOVNOSSVQVLLFPASVHAHAALFNYSISSRLSPRYWAS 300
QY      301  EAMLTSDLVMLGPGMAQMGTVLGFLORGAQLHEFPQYVKTALATDPAFCGALGEREG 360
DB      301  EAMLTSDLVMLGPGMAQMGTVLGFLORGAQLHEFPQYVKTALATDPAFCGALGEREG 360
QY      361  LEEDVVGORCPQDCITLQNVSAGLNHQTFSVYAAVSVAAQLNHTLQCNASGCPADDP 420
DB      361  LEEDVVGORCPQDCITLQNVSAGLNHQTFSVYAAVSVAAQLNHTLQCNASGCPADDP 420
QY      421  VAPWOLLEMMVLTFTVVGSLPLRPDSGQVNDMEYDLKLVMMWQSGVPRLLHVDGRFNGSLRT 480
DB      421  VAPWOLLEMMVLTFTVVGSLPLRPDSGQVNDMEYDLKLVMMWQSGVPRLLHVDGRFNGSLRT 480
QY      481  ERLKTRMTSDNQKQEVSRCSROCOBQVRRVKGFGHSCCYDCVDCBAGSYRQNPDIACTF 540
DB      481  ERLKTRMTSDNQKQEVSRCSROCOBQVRRVKGFGHSCCYDCVDCBAGSYRQNPDIACTF 540
QY      541  CGQDWSPEPRSTRCRRRRSRFLAMGEPAYLLLLLISLALGLVLAALGLFVHHRDSPLVQ 600
DB      541  CGQDWSPEPRSTRCRRRRSRFLAMGEPAYLLLLLISLALGLVLAALGLFVHHRDSPLVQ 600
QY      601  ASGGPLACFGVLCVGLSVLLFPQGPSPARCLAQOPLSHPLPTGCGSLTFLQAAEFIV 660
DB      601  ASGGPLACFGVLCVGLSVLLFPQGPSPARCLAQOPLSHPLPTGCGSLTFLQAAEFIV 660
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DB      601  ASGGPLACFGVLCVGLSVLLFPQGPSPARCLAQOPLSHPLPTGCGSLTFLQAAEFIV 660
QY      661  ESELPLSWADRLSGCLGPGPAMLVLLAMLVVALCTMYLVAFPEPVTTDMHMLPTETLV 720
DB      661  ESELPLSWADRLSGCLGPGPAMLVLLAMLVVALCTMYLVAFPEPVTTDMHMLPTETLV 720
QY      721  HCRTRSWVSGPLAATNATTAFLCFLGFLVRSQPGCYNRARGLTFAMLAFTVWSFVP 780
DB      721  HCRTRSWVSGPLAATNATTAFLCFLGFLVRSQPGCYNRARGLTFAMLAFTVWSFVP 780
QY      781  LLANVOVLRPAVQMGALLLCVGLIILAFHLPRCYLLMRQGLNTPPEFLGSGGPDAGQ 840
DB      781  LLANVOVLRPAVQMGALLLCVGLIILAFHLPRCYLLMRQGLNTPPEFLGSGGPDAGQ 840
QY      841  NDGNTGNQKHE 852
DB      841  NDGNTGNQKHE 852
```

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RESULT 7
US-09-796-338A-14
; Sequence 14, Application US/09796338A
; Patent No. US20020061522A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; TITLE OF INVENTION: 1983, 52881, 2398, 45449, 50289, AND
; TITLE OF INVENTION: 52872, NOVEL G PROTEIN-COUPLED RECEPTORS AND USES THEREFOR
; FILE REFERENCE: 10448-020001
; CURRENT FILING DATE: US/09/796, 338A
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: US 60/186, 059
; PRIOR FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 26
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-796-338A-14
```

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Query Match      99.7%; Score 4512; DB 9; Length 852;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1  MCGPAVLGLSLMALHPGTGAPLCLISQQLRMKGDIYVGLFPLGAEAEAGLRSRTRPSSP 60
DB      1  MCGPAVLGLSLMALHPGTGAPLCLISQQLRMKGDIYVGLFPLGAEAEAGLRSRTRPSSP 60
QY      61  VCTRSSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFPTCSEPPVAMKPSLMFLAKA 120
DB      61  VCTRSSNGLLMALMKAAVEEINNKSDLLPGLRLGYDLFPTCSEPPVAMKPSLMFLAKA 120
QY      121  GSRDIAAYCNTYOQPRVLAVIGPHSSBLAVTGFKEFSPFLMPQVSYGASMEILSARETF 180
DB      121  GSRDIAAYCNTYOQPRVLAVIGPHSSBLAVTGFKEFSPFLMPQVSYGASMEILSARETF 180
QY      181  PSFFPTVSDRYQLTAAAEELLQEFGMNVVAALGSDDEYGRQGLSTFSALAAARGICIAHE 240
DB      181  PSFFPTVSDRYQLTAAAEELLQEFGMNVVAALGSDDEYGRQGLSTFSALAAARGICIAHE 240
QY      241  GLVPLPRADDSRLGKVQDVLLHOVNOSSVQVLLFPASVHAHAALFNYSISSRLSPRYWAS 300
DB      241  GLVPLPRADDSRLGKVQDVLLHOVNOSSVQVLLFPASVHAHAALFNYSISSRLSPRYWAS 300
QY      301  EAMLTSDLVMLGPGMAQMGTVLGFLORGAQLHEFPQYVKTALATDPAFCGALGEREG 360
DB      301  EAMLTSDLVMLGPGMAQMGTVLGFLORGAQLHEFPQYVKTALATDPAFCGALGEREG 360
QY      361  LEEDVVGORCPQDCITLQNVSAGLNHQTFSVYAAVSVAAQLNHTLQCNASGCPADDP 420
DB      361  LEEDVVGORCPQDCITLQNVSAGLNHQTFSVYAAVSVAAQLNHTLQCNASGCPADDP 420
```

Qy 421 VKPQQLLENNYNTLFHHVGLPLRFDSSGNVDMEDYDLKLMWQGSVPRLHDVGRNGSLRT 480
Db 421 VKPQQLLENNYNTLFHHVGLPLRFDSSGNVDMEDYDLKLMWQGSVPRLHDVGRNGSLRT 480
Qy 481 ERLKIRMTSDNQKPVSRCSRCQCGOVRVYKGFHSCCYDCVDCAGSYRONPDDIACTF 540
Db 481 ERLKIRMTSDNQKPVSRCSRCQCGOVRVYKGFHSCCYDCVDCAGSYRONPDDIACTF 540
Qy 541 CGODEWSPERSTRCFRRRSRFLAWGEPAYLLLLLSLALGLVLAALGLFVHHRDSPLVQ 600
Db 541 CGODEWSPERSTRCFRRRSRFLAWGEPAYLLLLLSLALGLVLAALGLFVHHRDSPLVQ 600
Qy 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTFLQAAEIFV 660
Db 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTFLQAAEIFV 660
Qy 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAFPEVVTDMHMLPTEALV 720
Db 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAFPEVVTDMHMLPTEALV 720
Qy 721 HCRTRSWVSFGLAHATNATLAFCLCTGFLVRSQPGCYNRARGLTFAMLAFFITWVSFVP 780
Db 721 HCRTRSWVSFGLAHATNATLAFCLCTGFLVRSQPGCYNRARGLTFAMLAFFITWVSFVP 780
Qy 781 LLANVOVVLPAVQMGALLLCVGLIAAFHLPRCYLLMRQPGMLNTPPEFLGGGPGDAQOQ 840
Db 781 LLANVOVVLPAVQMGALLLCVGLIAAFHLPRCYLLMRQPGMLNTPPEFLGGGPGDAQOQ 840
Qy 841 NDGNTNGQKHE 852
Db 841 NDGNTNGQKHE 852

RESULT 8
US-10-261-482-2
; Sequence 2, Application US/10261482
; Publication No. US20030036089A1
; GENERAL INFORMATION:
; APPLICANT: WEI, Ming-Hui et al
; TITLE OF INVENTION: ISOLATED HUMAN G-PROTEIN COUPLED
; TITLE OF INVENTION: RECEPTORS, NUCLEIC ACID MOLECULES ENCODING HUMAN GPCR
; FILE REFERENCE: CLO00869CON
; CURRENT APPLICATION NUMBER: US/10/261,482
; PRIOR FILING DATE: 2002-10-02
; PRIOR APPLICATION NUMBER: 09/684,393
; PRIOR FILING DATE: 2000-10-10
; PRIOR APPLICATION NUMBER: 60/172,600
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 852
; TYPE: PRT
; ORGANISM: HUMAN
US-10-261-482-2

Query Match 99.7%; Score 4512; DB 14; Length 852;

Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 MLGPAVLGLSLWALLHPGTGAPLCLISQQLRMKGVDVVLGGLPPLGAEBAEAGLSRTRPSP 60
Db 1 MLGPAVLGLSLWALLHPGTGAPLCLISQQLRMKGVDVVLGGLPPLGAEBAEAGLSRTRPSP 60
Qy 61 VCTRSSNGGLMALAKMAVEEINKSDLLPGLRIGYDLPTDCSPFPVAMKPSLMFLAKA 120
Db 61 VCTRSSNGGLMALAKMAVEEINKSDLLPGLRIGYDLPTDCSPFPVAMKPSLMFLAKA 120
Qy 121 GSRDIAACNTQYQPRVLAVIGPHSSELAWTGKFSFPLMPOVSYGASMEILSARPTF 180
Db 121 GSRDIAACNTQYQPRVLAVIGPHSSELAWTGKFSFPLMPOVSYGASMEILSARPTF 180

Qy 181 PSFPRVPSDRVQJTAABELLQEFGMWVVAALSGDDEYRGQSLISFALAAARGICIAHE 240
Db 181 PSFPRVPSDRVQJTAABELLQEFGMWVVAALSGDDEYRGQSLISFALAAARGICIAHE 240
Qy 241 GLVPLPRADDSRLGKQVDVLHQNQSSVQVYVLLPASVHAHAALFNYSISSRLSPKMWAS 300
Db 241 GLVPLPRADDSRLGKQVDVLHQNQSSVQVYVLLPASVHAHAALFNYSISSRLSPKMWAS 300
Qy 301 EAMTSDLVNGLPGMAQMGTVLGFLOGAQLHEFPQYVKTHTALATPAFCALGEREOG 360
Db 301 EAMTSDLVNGLPGMAQMGTVLGFLOGAQLHEFPQYVKTHTALATPAFCALGEREOG 360
Qy 361 LEEDVVGQRCPCDCITLQNVASAGLNHRQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
Db 361 LEEDVVGQRCPCDCITLQNVASAGLNHRQTFSSVYAAYVSAQALHNTLQCNASGCPADP 420
Qy 421 VKPQQLLENNYNTLFHHVGLPLRFDSSGNVDMEDYDLKLMWQGSVPRLHDVGRNGSLRT 480
Db 421 VKPQQLLENNYNTLFHHVGLPLRFDSSGNVDMEDYDLKLMWQGSVPRLHDVGRNGSLRT 480
Qy 481 ERLKIRMTSDNQKPVSRCSRCQCGOVRVYKGFHSCCYDCVDCAGSYRONPDDIACTF 540
Db 481 ERLKIRMTSDNQKPVSRCSRCQCGOVRVYKGFHSCCYDCVDCAGSYRONPDDIACTF 540
Qy 541 CGODEWSPERSTRCFRRRSRFLAWGEPAYLLLLLSLALGLVLAALGLFVHHRDSPLVQ 600
Db 541 CGODEWSPERSTRCFRRRSRFLAWGEPAYLLLLLSLALGLVLAALGLFVHHRDSPLVQ 600
Qy 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTFLQAAEIFV 660
Db 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTFLQAAEIFV 660
Qy 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAFPEVVTDMHMLPTEALV 720
Db 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLVAFPEVVTDMHMLPTEALV 720
Qy 721 HCRTRSWVSFGLAHATNATLAFCLCTGFLVRSQPGCYNRARGLTFAMLAFFITWVSFVP 780
Db 721 HCRTRSWVSFGLAHATNATLAFCLCTGFLVRSQPGCYNRARGLTFAMLAFFITWVSFVP 780
Qy 781 LLANVOVVLPAVQMGALLLCVGLIAAFHLPRCYLLMRQPGMLNTPPEFLGGGPGDAQOQ 840
Db 781 LLANVOVVLPAVQMGALLLCVGLIAAFHLPRCYLLMRQPGMLNTPPEFLGGGPGDAQOQ 840
Qy 841 NDGNTNGQKHE 852
Db 841 NDGNTNGQKHE 852

RESULT 9
US-10-282-837-14
; Sequence 14, Application US/10282837
; Publication No. US20030082738A1
; GENERAL INFORMATION:
; APPLICANT: Millennium Pharmaceuticals, Inc.
; TITLE OF INVENTION: 52872, NOVEL G PROTEIN-COUPLED RECEPTORS AND USES THEREFOR
; FILE REFERENCE: 10448-020001
; CURRENT APPLICATION NUMBER: US/10/282,837
; PRIOR FILING DATE: 2002-10-29
; PRIOR APPLICATION NUMBER: US/09/796,338
; PRIOR FILING DATE: 2001-02-28
; PRIOR APPLICATION NUMBER: US 60/186,059
; PRIOR FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 26
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-282-837-14

Query Match 99.7%; Score 4512; DB 14; Length 852;

Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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QY 1 MGPAVLGSLWALHPGTGAPLCLSQQLRMKGDYVLCGLFPLGEAEAGLSRTRPSSP 60
DB 1 MUGPAVLGSLWALHPGTGAPLCLSQQLRMKGDYVLCGLFPLGEAEAGLSRTRPSSP 60
QY 61 VCTRSSNGLLMALMKMAVEEINNKSDDLPLGLRGLGYDLPDTCSEPVVAMKPSLMLFLAKA 120
DB 61 VCTRSSNGLLMALMKMAVEEINNKSDDLPLGLRGLGYDLPDTCSEPVVAMKPSLMLFLAKA 120
QY 121 GSRDIAAYCNTYOQPRVLAVIGPHSSSELAMVTGKFFSFLMPQVSYGASMEILLARETF 180
DB 121 GSRDIAAYCNTYOQPRVLAVIGPHSSSELAMVTGKFFSFLMPQVSYGASMEILLARETF 180
QY 181 PSFPRVPSDRVQLTAAAEILQEFQGNWVAALGSDDEYRGQGLSIFSAALAAAGICIAHE 240
DB 181 PSFPRVPSDRVQLTAAAEILQEFQGNWVAALGSDDEYRGQGLSIFSAALAAAGICIAHE 240
QY 241 GLVPLPRADDSRLGKVQDVLHQVNOSSVOVVLFPASVHAHALFNYSISSRLSPKVMVAS 300
DB 241 GLVPLPRADDSRLGKVQDVLHQVNOSSVOVVLFPASVHAHALFNYSISSRLSPKVMVAS 300
QY 301 EAMLTSDLVNGLPGMAQMGTVLGLQGAQLHEFPQYKTHALATDPAFCALGEREG 360
DB 301 EAMLTSDLVNGLPGMAQMGTVLGLQGAQLHEFPQYKTHALATDPAFCALGEREG 360
QY 361 LEBDVVQRCPCQDCITLQNVSAAGLNHQTFSYVAAYVSAQALHNTLQCNASGCPADP 420
DB 361 LEBDVVQRCPCQDCITLQNVSAAGLNHQTFSYVAAYVSAQALHNTLQCNASGCPADP 420
QY 421 VAPMOLLENNYLTFFHVGGLPLRFDSSGNVDMEDYDKLWMOGSAVRLHDVGRFNGSLRT 480
DB 421 VAPMOLLENNYLTFFHVGGLPLRFDSSGNVDMEDYDKLWMOGSAVRLHDVGRFNGSLRT 480
QY 481 ERLKIRMTSDNQKPVSRSCROCEGOVARVKGFSHCYDCCVDCBAGSYRONDDIACF 540
DB 481 ERLKIRMTSDNQKPVSRSCROCEGOVARVKGFSHCYDCCVDCBAGSYRONDDIACF 540
QY 541 CGODEMSPERSTRCFRRSRFLANGEPAYLLLLLSLALGLVLAALGLFVHHRSPLVQ 600
DB 541 CGODEMSPERSTRCFRRSRFLANGEPAYLLLLLSLALGLVLAALGLFVHHRSPLVQ 600
QY 601 ASGGPLACGVLGVLGVLCLSVLLPFGQSPARCIAQOPLSHLPLTGCLSTLFLQAAELFV 660
DB 601 ASGGPLACGVLGVLGVLCLSVLLPFGQSPARCIAQOPLSHLPLTGCLSTLFLQAAELFV 660
QY 661 ESELPLSWADRLSGCLRGPMAMLVLLAMLEVALCTMYLVAFPPEVVTDMHMLPTEALV 720
DB 661 ESELPLSWADRLSGCLRGPMAMLVLLAMLEVALCTMYLVAFPPEVVTDMHMLPTEALV 720
QY 721 HCRTRSWVSFGLAHATNATLAFCLGTLFVRSQPCYNRARGLTFAMLAAYITWVSFVP 780
DB 721 HCRTRSWVSFGLAHATNATLAFCLGTLFVRSQPCYNRARGLTFAMLAAYITWVSFVP 780
QY 781 LIANVOVLRPAVONGALLCYLGLIAAHLRCTLAMPQGLNTPPEFLGGGPDAAQO 840
DB 781 LIANVOVLRPAVONGALLCYLGLIAAHLRCTLAMPQGLNTPPEFLGGGPDAAQO 840
QY 841 NDGNTGNOGKHE 852
DB 841 NDGNTGNOGKHE 852

```

RESULT 10
US-10-145-586-14

; Sequence 14, Application US/10145586
; Publication No. US20030138890A1
; GENERAL INFORMATION:
; APPLICANT: Alexandra Glucksmann, Maria
; APPLICANT: Silvio-Santiago, Immaculada
; APPLICANT: M. Galvin, Katherine
; APPLICANT: Weich, Nadine

```

; APPLICANT: Curtis, Rory A.J.
; APPLICANT: Bandaru, Rajasekhar
; APPLICANT: Kapeller-Lieberman, Rosana
; TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTOR FAMILY MEMBERS,
; TITLE OF INVENTION: HUMAN THIOREDOXIN FAMILY MEMBERS, HUMAN LEUCINE-RICH
; TITLE OF INVENTION: REPEAT FAMILY MEMBERS, AND HUMAN RINGFINGER FAMILY MEMBER
; FILE REFERENCE: 10448-188001
; CURRENT APPLICATION NUMBER: US/10/145,586
; CURRENT FILING DATE: 2002-05-14
; NUMBER OF SEQ ID NOS: 95
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 852
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-145-586-14

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Query Match 99.7%; Score 4512; DB 14; Length 852;

Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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QY 1 MUGPAVLGSLWALHPGTGAPLCLSQQLRMKGDYVLCGLFPLGEAEAGLSRTRPSSP 60
DB 1 MUGPAVLGSLWALHPGTGAPLCLSQQLRMKGDYVLCGLFPLGEAEAGLSRTRPSSP 60
QY 61 VCTRSSNGLLMALMKMAVEEINNKSDDLPLGLRGLGYDLPDTCSEPVVAMKPSLMLFLAKA 120
DB 61 VCTRSSNGLLMALMKMAVEEINNKSDDLPLGLRGLGYDLPDTCSEPVVAMKPSLMLFLAKA 120
QY 121 GSRDIAAYCNTYOQPRVLAVIGPHSSSELAMVTGKFFSFLMPQVSYGASMEILLARETF 180
DB 121 GSRDIAAYCNTYOQPRVLAVIGPHSSSELAMVTGKFFSFLMPQVSYGASMEILLARETF 180
QY 181 PSFPRVPSDRVQLTAAAEILQEFQGNWVAALGSDDEYRGQGLSIFSAALAAAGICIAHE 240
DB 181 PSFPRVPSDRVQLTAAAEILQEFQGNWVAALGSDDEYRGQGLSIFSAALAAAGICIAHE 240
QY 241 GLVPLPRADDSRLGKVQDVLHQVNOSSVOVVLFPASVHAHALFNYSISSRLSPKVMVAS 300
DB 241 GLVPLPRADDSRLGKVQDVLHQVNOSSVOVVLFPASVHAHALFNYSISSRLSPKVMVAS 300
QY 301 EAMLTSDLVNGLPGMAQMGTVLGLQGAQLHEFPQYKTHALATDPAFCALGEREG 360
DB 301 EAMLTSDLVNGLPGMAQMGTVLGLQGAQLHEFPQYKTHALATDPAFCALGEREG 360
QY 361 LEBDVVQRCPCQDCITLQNVSAAGLNHQTFSYVAAYVSAQALHNTLQCNASGCPADP 420
DB 361 LEBDVVQRCPCQDCITLQNVSAAGLNHQTFSYVAAYVSAQALHNTLQCNASGCPADP 420
QY 421 VAPMOLLENNYLTFFHVGGLPLRFDSSGNVDMEDYDKLWMOGSAVRLHDVGRFNGSLRT 480
DB 421 VAPMOLLENNYLTFFHVGGLPLRFDSSGNVDMEDYDKLWMOGSAVRLHDVGRFNGSLRT 480
QY 481 ERLKIRMTSDNQKPVSRSCROCEGOVARVKGFSHCYDCCVDCBAGSYRONDDIACF 540
DB 481 ERLKIRMTSDNQKPVSRSCROCEGOVARVKGFSHCYDCCVDCBAGSYRONDDIACF 540
QY 541 CGODEMSPERSTRCFRRSRFLANGEPAYLLLLLSLALGLVLAALGLFVHHRSPLVQ 600
DB 541 CGODEMSPERSTRCFRRSRFLANGEPAYLLLLLSLALGLVLAALGLFVHHRSPLVQ 600
QY 601 ASGGPLACGVLGVLGVLCLSVLLPFGQSPARCIAQOPLSHLPLTGCLSTLFLQAAELFV 660
DB 601 ASGGPLACGVLGVLGVLCLSVLLPFGQSPARCIAQOPLSHLPLTGCLSTLFLQAAELFV 660
QY 661 ESELPLSWADRLSGCLRGPMAMLVLLAMLEVALCTMYLVAFPPEVVTDMHMLPTEALV 720
DB 661 ESELPLSWADRLSGCLRGPMAMLVLLAMLEVALCTMYLVAFPPEVVTDMHMLPTEALV 720
QY 721 HCRTRSWVSFGLAHATNATLAFCLGTLFVRSQPCYNRARGLTFAMLAAYITWVSFVP 780
DB 721 HCRTRSWVSFGLAHATNATLAFCLGTLFVRSQPCYNRARGLTFAMLAAYITWVSFVP 780

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Qy 761 LLANVOVVLPRPVGKALLLCVLTGILAAFHLPKCYLMMKROPLNTPPEFFLGCGPGDAGQ 840
Db 761 LLANVOVVLPRPVGKALLLCVLTGILAAFHLPKCYLMMKROPLNTPPEFFLGCGPGDAGQ 840
Qy 841 NDGNTGNGKHE 852
Db 841 NDGNTGNGKHE 852

RESULT 11
US-10-407-079-90
Sequence 90, Application US/10407079
Publication No. US20030215860A1
GENERAL INFORMATION:
APPLICANT: Millennium Pharmaceuticals, Inc.
APPLICANT: Gluckmann, Maria A.
APPLICANT: Silos-Santiago, Immaculada
APPLICANT: Carroll, Joseph M.
APPLICANT: Galvin, Katherine M.
TITLE OF INVENTION: 16636, 2466, 43238, 1983, 52881, 2398,
TITLE OF INVENTION: 45449, 50289, 52872 AND 26908 MOLECULES AND USES THEREFOR
FILE REFERENCE: MP103-0510MNTM
CURRENT FILING DATE: 2003-04-03
PRIOR APPLICATION NUMBER: US/10/407,079
PRIOR FILING DATE: 2002-08-22
PRIOR APPLICATION NUMBER: US 60/314,041
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: US 10/225,094
PRIOR FILING DATE: 2002-08-21
PRIOR APPLICATION NUMBER: US 60/314,185
PRIOR FILING DATE: 2001-08-22
PRIOR APPLICATION NUMBER: US 10/272,417
PRIOR FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: US 09/715,790
PRIOR FILING DATE: 2000-11-17
PRIOR APPLICATION NUMBER: US 60/191,845
PRIOR FILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: US 10/282,837
PRIOR FILING DATE: 2002-10-29
PRIOR APPLICATION NUMBER: US 09/796,338
PRIOR FILING DATE: 2001-02-28
PRIOR APPLICATION NUMBER: US 60/186,059
PRIOR FILING DATE: 2000-02-29
Remaining Prior Application data removed - See file wrapper or PALM.
NUMBER OF SEQ ID NOS: 110
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 90
LENGTH: 852
TYPE: PRT
ORGANISM: Homo sapiens
US-10-407-079-90

Query Match 99.7%; Score 4512; DB 15; Length 852;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 241 GLVPLPRADSRRLGKQVDVILHQVNOSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
Db 241 GLVPLPRADSRRLGKQVDVILHQVNOSSVQVVLFPASVHAHALFNYSISSRLSPKVVAS 300
Qy 301 EAWLTSDLVNGLPMAQMGTVLGFLOQAOHLHEFPQYKTHLATDPAFCSALGERBOG 360
Db 301 EAWLTSDLVNGLPMAQMGTVLGFLOQAOHLHEFPQYKTHLATDPAFCSALGERBOG 360
Qy 361 LEEDVNGORCPQCCITLQVNSAGLNHOFPSVYAAYVAQALHNTLQCNASGCPNDP 420
Db 361 LEEDVNGORCPQCCITLQVNSAGLNHOFPSVYAAYVAQALHNTLQCNASGCPNDP 420
Qy 421 VKPQQLLENNYNTLFHVGGLPLRFDSGGNDMEYDLKLWQGSVPLHVDGRFNGSLRT 480
Db 421 VKPQQLLENNYNTLFHVGGLPLRFDSGGNDMEYDLKLWQGSVPLHVDGRFNGSLRT 480
Qy 481 ERLKIRWHTSDNQKRVSCRSQCEGQVRRVKGHSCCYDCVDCAGSYRONPDIACTF 540
Db 481 ERLKIRWHTSDNQKRVSCRSQCEGQVRRVKGHSCCYDCVDCAGSYRONPDIACTF 540
Qy 541 CGQDENSPESTRCFRRRSRFLAMGEPAYVLLLLLSALGVLAALGLFVHHRDSPVQ 600
Db 541 CGQDENSPESTRCFRRRSRFLAMGEPAYVLLLLLSALGVLAALGLFVHHRDSPVQ 600
Qy 601 ASGGPLACFGVLCGLVCLSVLFPQGPSPARCIAQOPLSLPTGCLSTFLQAAEIFV 660
Db 601 ASGGPLACFGVLCGLVCLSVLFPQGPSPARCIAQOPLSLPTGCLSTFLQAAEIFV 660
Qy 661 ESELPLSMADRLSGCLGPGPAMVLVLLAMLEVALCTWYLVAPPEVVTDMHMLPTBALV 720
Db 661 ESELPLSMADRLSGCLGPGPAMVLVLLAMLEVALCTWYLVAPPEVVTDMHMLPTBALV 720
Qy 721 HCRTRSWVSFGLAATNATLAFCLGTPLVRSGPGVYNAARGTFLMAYFTWVSFVP 780
Db 721 HCRTRSWVSFGLAATNATLAFCLGTPLVRSGPGVYNAARGTFLMAYFTWVSFVP 780
Qy 781 LLANVOVVLPRPVGKALLLCVLTGILAAFHLPKCYLMMKROPLNTPPEFFLGCGPGDAGQ 840
Db 781 LLANVOVVLPRPVGKALLLCVLTGILAAFHLPKCYLMMKROPLNTPPEFFLGCGPGDAGQ 840
Qy 841 NDGNTGNGKHE 852
Db 841 NDGNTGNGKHE 852

RESULT 12
US-10-179-373-7
Sequence 7, Application US/10179373
Publication No. US20030232407A1
GENERAL INFORMATION:
APPLICANT: ZOLLER, MARK
APPLICANT: LI, XIAODONG
APPLICANT: STASZEWSKI, LENA
APPLICANT: O'CONNELL, SHAWN
APPLICANT: ZOCULYA, SERGEY
APPLICANT: ADLER, JON
APPLICANT: XU, HONG
APPLICANT: ECHEVERRI, FERNANDO
TITLE OF INVENTION: T1R HETERO-OLIGOMERIC TASTE RECEPTORS AND CELL LINES
TITLE OF INVENTION: THAT EXPRESS SAID RECEPTORS AND USE THEREOF FOR
FILE REFERENCE: 078003-0291566
CURRENT APPLICATION NUMBER: US/10/179,373
CURRENT FILING DATE: 2002-06-26
PRIOR APPLICATION NUMBER: 60/300,434
PRIOR FILING DATE: 2001-06-26
PRIOR APPLICATION NUMBER: 60/304,749
PRIOR FILING DATE: 2001-07-13
PRIOR APPLICATION NUMBER: 60/310,493
PRIOR FILING DATE: 2001-08-08
PRIOR APPLICATION NUMBER: 60/331,771
PRIOR FILING DATE: 2001-11-21

; PRIOR APPLICATION NUMBER: 60/339,472
 ; PRIOR FILING DATE: 2001-12-14
 ; PRIOR APPLICATION NUMBER: 60/372,090
 ; PRIOR FILING DATE: 2002-04-15
 ; PRIOR APPLICATION NUMBER: 60/374,143
 ; PRIOR FILING DATE: 2002-04-22
 ; NUMBER OF SEQ ID NOS: 19
 ; SOFTWARE: Patent In Ver. 2.1
 ; SEQ ID NO 7
 ; LENGTH: 852
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-179-373-7

Query Match 99.7%; Score 4512; DB 15; Length 852;
 Best Local Similarity 99.9%; Pred. No. 0;
 Matches 851; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 MGPAVLGLSLWALLHPGTGAPLCLSQQLRMKGIVLGGFLPGEAEAGLRSTRPSSP 60
 DB 1 MGPAVLGLSLWALLHPGTGAPLCLSQQLRMKGIVLGGFLPGEAEAGLRSTRPSSP 60
 QY 61 VCTRSSNGLLMALMKAAVEEINKSDLLPGLRLGYDLPTCSEPVVAMKPSLMFLAKA 120
 DB 61 VCTRSSNGLLMALMKAAVEEINKSDLLPGLRLGYDLPTCSEPVVAMKPSLMFLAKA 120
 QY 121 GSRDIAAYCNTYOQPRVLAIVGPHSSSLAWTGKFFSFLMPOVSYGASWELLSARETF 180
 DB 121 GSRDIAAYCNTYOQPRVLAIVGPHSSSLAWTGKFFSFLMPOVSYGASWELLSARETF 180
 QY 181 PSFRTVSDRVQVLTAAELLQEFGMNVAALGSDDEYGRQGLSIFSALAAARGICIAHE 240
 DB 181 PSFRTVSDRVQVLTAAELLQEFGMNVAALGSDDEYGRQGLSIFSALAAARGICIAHE 240
 QY 241 GLVPLPRADDSRLGKVQDVHLHQVNSVQVLLFASVHAHALFNYSSISRLSPKVVAS 300
 DB 241 GLVPLPRADDSRLGKVQDVHLHQVNSVQVLLFASVHAHALFNYSSISRLSPKVVAS 300
 QY 301 EAMLTSDLVWGLPGMAQGTVLGFLQGAQLHEFPQVYKTHALATDPAFCGALGERBOG 360
 DB 301 EAMLTSDLVWGLPGMAQGTVLGFLQGAQLHEFPQVYKTHALATDPAFCGALGERBOG 360
 QY 361 LEEDVQGRCPQCCITLQNVSAAGLNHHQTFSVYAAYSVAQALNHTLQCNASGCPADP 420
 DB 361 LEEDVQGRCPQCCITLQNVSAAGLNHHQTFSVYAAYSVAQALNHTLQCNASGCPADP 420
 QY 421 VKPMQLLENNYNTLTFHVGGPLRFDSSGNVMEYDLKLMWQGSVPRLHDVGRFNGSLRT 480
 DB 421 VKPMQLLENNYNTLTFHVGGPLRFDSSGNVMEYDLKLMWQGSVPRLHDVGRFNGSLRT 480
 QY 481 ERKTRMHTSDNOKVSRCSROCGOVRVKGPHSSCCYDCYDCENAGSRONPDIACTF 540
 DB 481 ERKTRMHTSDNOKVSRCSROCGOVRVKGPHSSCCYDCYDCENAGSRONPDIACTF 540
 QY 541 CGODEMSEPSRSTRCRRRSRFLAMGEPAYLILLLLSLALGLVLAALGLFVHHRDSPVQ 600
 DB 541 CGODEMSEPSRSTRCRRRSRFLAMGEPAYLILLLLSLALGLVLAALGLFVHHRDSPVQ 600
 QY 601 ASGGLACFGIVCLGVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTLFLQAAELFV 660
 DB 601 ASGGLACFGIVCLGVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTLFLQAAELFV 660
 QY 661 ESELPLSWADRLSGCLRGPMWLVVLLMLVEVALCTWYLVAFPEVVYTDWMTLEALV 720
 DB 661 ESELPLSWADRLSGCLRGPMWLVVLLMLVEVALCTWYLVAFPEVVYTDWMTLEALV 720
 QY 721 HCRTRSWVSFGLAHATNATLAFCLGFTLVRSOPGVNBARGLFAMLAAYITWVSFP 780
 DB 721 HCRTRSWVSFGLAHATNATLAFCLGFTLVRSOPGVNBARGLFAMLAAYITWVSFP 780
 QY 781 LLANQVVLPRAVQWGAALLCVLGLILAAFLPRCYLWKQPGLANTPPEFLGGGPGDAQO 840
 DB 781 LLANQVVLPRAVQWGAALLCVLGLILAAFLPRCYLWKQPGLANTPPEFLGGGPGDAQO 840

QY 841 NDGNTNGCKHE 852
 DB 841 NDGNTNGCKHE 852

RESULT 13
 US-10-280-183A-5
 ; Publication 5, Application US/10280183A
 ; Publication No. US20040081964A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Pfizer Inc.
 ; APPLICANT: Bachmanov, Alexander A
 ; APPLICANT: Beauchamp, Gary K.
 ; APPLICANT: Chatterjee, Anubindo
 ; APPLICANT: De Jong, Pieter J.
 ; APPLICANT: Li, Shanru
 ; APPLICANT: Li, Xia
 ; APPLICANT: Ohmen, Jeffrey D
 ; APPLICANT: Reed, Danielle R.
 ; APPLICANT: Ross, David
 ; APPLICANT: Tordoff, Michael G.
 ; TITLE OF INVENTION: GENE AND SEQUENCE VARIATION ASSOCIATED WITH SENSING
 ; TITLE OF INVENTION: CARBOHYDRATE COMPOUNDS AND OTHER SWEETENERS
 ; FILE REFERENCE: PCI8306A
 ; CURRENT APPLICATION NUMBER: US/10/280,183A
 ; PRIOR APPLICATION NUMBER: 60/200,794
 ; PRIOR FILING DATE: 2000-04-28
 ; NUMBER OF SEQ ID NOS: 652
 ; SOFTWARE: Patent In Ver. 3.1
 ; SEQ ID NO 5
 ; LENGTH: 852
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-280-183A-5

Query Match 99.7%; Score 4509; DB 16; Length 852;
 Best Local Similarity 99.8%; Pred. No. 0;
 Matches 850; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 MGPAVLGLSLWALLHPGTGAPLCLSQQLRMKGIVLGGFLPGEAEAGLRSTRPSSP 60
 DB 1 MGPAVLGLSLWALLHPGTGAPLCLSQQLRMKGIVLGGFLPGEAEAGLRSTRPSSP 60
 QY 61 VCTRSSNGLLMALMKAAVEEINKSDLLPGLRLGYDLPTCSEPVVAMKPSLMFLAKA 120
 DB 61 VCTRSSNGLLMALMKAAVEEINKSDLLPGLRLGYDLPTCSEPVVAMKPSLMFLAKA 120
 QY 121 GSRDIAAYCNTYOQPRVLAIVGPHSSSLAWTGKFFSFLMPOVSYGASWELLSARETF 180
 DB 121 GSRDIAAYCNTYOQPRVLAIVGPHSSSLAWTGKFFSFLMPOVSYGASWELLSARETF 180
 QY 181 PSFRTVSDRVQVLTAAELLQEFGMNVAALGSDDEYGRQGLSIFSALAAARGICIAHE 240
 DB 181 PSFRTVSDRVQVLTAAELLQEFGMNVAALGSDDEYGRQGLSIFSALAAARGICIAHE 240
 QY 241 GLVPLPRADDSRLGKVQDVHLHQVNSVQVLLFASVHAHALFNYSSISRLSPKVVAS 300
 DB 241 GLVPLPRADDSRLGKVQDVHLHQVNSVQVLLFASVHAHALFNYSSISRLSPKVVAS 300
 QY 301 EAMLTSDLVWGLPGMAQGTVLGFLQGAQLHEFPQVYKTHALATDPAFCGALGERBOG 360
 DB 301 EAMLTSDLVWGLPGMAQGTVLGFLQGAQLHEFPQVYKTHALATDPAFCGALGERBOG 360
 QY 361 LEEDVQGRCPQCCITLQNVSAAGLNHHQTFSVYAAYSVAQALNHTLQCNASGCPADP 420
 DB 361 LEEDVQGRCPQCCITLQNVSAAGLNHHQTFSVYAAYSVAQALNHTLQCNASGCPADP 420
 QY 421 VKPMQLLENNYNTLTFHVGGPLRFDSSGNVMEYDLKLMWQGSVPRLHDVGRFNGSLRT 480
 DB 421 VKPMQLLENNYNTLTFHVGGPLRFDSSGNVMEYDLKLMWQGSVPRLHDVGRFNGSLRT 480

Qy 481 ERLKIRMTSDNOKPVSRCSCQCEGOVRRVKGFSHCYDCEVDCBAGSYRONPDDIACTF 540
Db 481 ERLKIRMTSDNOKPVSRCSCQCEGOVRRVKGFSHCYDCEVDCBAGSYRONPDDIACTF 540
Qy 541 CGODEMSPERSTRCFRRSRFLAMGEPAYLLILLSLALGLVLAALGLFVHHRDSPLVQ 600
Db 541 CGODEMSPERSTRCFRRSRFLAMGEPAYLLILLSLALGLVLAALGLFVHHRDSPLVQ 600
Qy 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTLFLQAAELFV 660
Db 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTLFLQAAELFV 660
Qy 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLAFPPEVVTDMHMLPTEALV 720
Db 661 ESELPLSMADRLSGCLRGPMAMLVLLMLVEVALCTWYLAFPPEVVTDMHMLPTEALV 720
Qy 721 HCRTRSWVSFGIAHATNATLAFLCFLGTFLVRSQGCYNRAGLTFPAMLAFFITWVSFVP 780
Db 721 HCRTRSWVSFGIAHATNATLAFLCFLGTFLVRSQGCYNRAGLTFPAMLAFFITWVSFVP 780
Qy 781 LLANQVVLPRPAVOMGALLLCVIGIILAAFHLPRCYLIMRQPGINTPEFFLGGGPDQAQO 840
Db 781 LLANQVVLPRPAVOMGALLLCVIGIILAAFHLPRCYLIMRQPGINTPEFFLGGGPDQAQO 840
Qy 841 NDGNTGNQKHE 852
Db 841 NDGNTGNQKHE 852

RESULT 14

us-09-799-629-4
; Sequence 4, Application US/09799629
; Publication No. US2003008344A1
; GENERAL INFORMATION:
; APPLICANT: ADLER, JON ELIOT
; APPLICANT: ZOZULYA, SERGEY
; APPLICANT: LI, XIADONG
; APPLICANT: O'CONNELL, SHAM
; APPLICANT: STASZEWSKI, LENA
; TITLE OF INVENTION: T1R TASTE RECEPTORS AND GENES ENCODING SAME
; FILE REFERENCE: 078003/027870/RXT
; CURRENT APPLICATION NUMBER: US/09/799,629
; PRIOR FILING DATE: 2001-03-07
; PRIOR APPLICATION NUMBER: 60/187,546
; PRIOR FILING DATE: 2000-03-07
; PRIOR APPLICATION NUMBER: 60/195,536
; PRIOR FILING DATE: 2000-04-07
; PRIOR APPLICATION NUMBER: 60/209,840
; PRIOR FILING DATE: 2000-06-06
; PRIOR APPLICATION NUMBER: 60/214,213
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: 60/226,448
; PRIOR FILING DATE: 2000-08-17
; PRIOR APPLICATION NUMBER: 60/259,227
; PRIOR FILING DATE: 2001-01-03
; NUMBER OF SEQ ID NOS: 20
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 850
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-799-629-4

Query Match 99.5%; Score 4500; DB 10; Length 850;
Best Local Similarity 99.6%; Pred. No. 0;
Matches 849; Conservative 0; Mismatches 1; Indels 2; Gaps 1;
Qy 1 MLGPVILSLIMALLHPGTGAAPLCLSQQLRMKGDVVLGGLPLGAEABAGLSRTRRSPSP 60
Db 1 MLGPVILSLIMALLHPGTGAAPLCLSQQLRMKGDVVLGGLPLGAEABAGLSRTRRSPSP 60
Qy 61 VCTRFSSNGLMALAMKAVEEINNKSDLLPGLRLGYDLFDPTCSEPVVAMKPSIMFLAKA 120
Db 61 VCTRFSSNGLMALAMKAVEEINNKSDLLPGLRLGYDLFDPTCSEPVVAMKPSIMFLAKA 120

Db 61 VCTRFSSNGLMALAMKAVEEINNKSDLLPGLRLGYDLFDPTCSEPVVAMKPSIMFLAKA 120
Qy 121 GSRDIAAYCNTYQYOPRLAVIYGHSSSELMAMVTGKFPSPFLMPQVSGASWELLSARETF 180
Db 121 GSRDIAAYCNTYQYOPRLAVIYGHSSSELMAMVTGKFPSPFLMPH--YGASWELLSARETF 178
Qy 181 PSFRTYPSDRVQYTAABELLQEFGMWVAALGSDDEYGRQGLSIFSALAAARGICIAHE 240
Db 179 PSFRTYPSDRVQYTAABELLQEFGMWVAALGSDDEYGRQGLSIFSALAAARGICIAHE 238
Qy 241 GLVPLPRADSRRLKQVDVHQNQSSVQVYLLPASVAAHALFNYSISSRLSPKVVAS 300
Db 239 GLVPLPRADSRRLKQVDVHQNQSSVQVYLLPASVAAHALFNYSISSRLSPKVVAS 298
Qy 301 EAMLTSPLVMGLPMAQMGTVLGFLOGAOLHEFPQYVKTHLALATPAFCALSAGEEOG 360
Db 299 EAMLTSPLVMGLPMAQMGTVLGFLOGAOLHEFPQYVKTHLALATPAFCALSAGEEOG 358
Qy 361 LEEBVGQRCPQDCITLQNVSAGLNHRQTFSSVYAAVYAQALHNTLQCNASGCPAODP 420
Db 359 LEEBVGQRCPQDCITLQNVSAGLNHRQTFSSVYAAVYAQALHNTLQCNASGCPAODP 418
Qy 421 VKPWQLLENNYNTLTFHYGGLPLRFDSGSQVMEYDLKLMWQGSVPRLHVGFRNGSLRT 480
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Qy 481 ERLKIRMTSDNOKPVSRCSCQCEGOVRRVKGFSHCYDCEVDCBAGSYRONPDDIACTF 540
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Qy 541 CGODEMSPERSTRCFRRSRFLAMGEPAYLLILLSLALGLVLAALGLFVHHRDSPLVQ 600
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Qy 601 ASGGPLACFGVLCGLVCLSVLLFPQGPSPARCLAQOPLSHLPLTGCLSTLFLQAAELFV 660
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Qy 841 NDGNTGNQKHE 852
Db 839 NDGNTGNQKHE 850

RESULT 15

US-10-190-417-30
; Sequence 30, Application US/10190417
; Publication No. US20030166137A1
; GENERAL INFORMATION:
; APPLICANT: Zuker, Charles S.
; APPLICANT: Ryba, Nicholas J. P.
; APPLICANT: Chandrasekar, Jayaram
; APPLICANT: Hoon, Mark A.
; APPLICANT: Nelson, Greg
; APPLICANT: Zhang, Yifeng
; APPLICANT: The Regents of the University of California
; APPLICANT: as represented by the Secretary of the
; APPLICANT: Department of Health and Human Services
; TITLE OF INVENTION: Mammalian Sweet and Amino Acid Heterodimeric Taste
; FILE REFERENCE: 02307E-120130US

1 CURRENT APPLICATION NUMBER: US/10/190,417
1 CURRENT FILING DATE: 2002-11-14
1 PRIOR APPLICATION NUMBER: US 60/302,898
1 PRIOR FILING DATE: 2001-07-03
1 PRIOR APPLICATION NUMBER: US 09/927,315
1 PRIOR FILING DATE: 2001-08-10
1 PRIOR APPLICATION NUMBER: US 60/358,925
1 PRIOR FILING DATE: 2002-02-22
1 NUMBER OF SEQ ID NOS: 30
1 SOFTWARE: PatentIn Ver. 2.1
1 SEQ ID NO: 30
1 LENGTH: 850
1 TYPE: PRT
1 ORGANISM: Homo sapiens
1 FEATURE:
1 OTHER INFORMATION: human TIR3
1 US-10-190-417-30

Query Match 99.5%; Score 4500; DB 14; Length 850;
Best Local Similarity 99.6%; Pred. No. 0;
Matches 849; Conservative 0; Mismatches 1; Indels 2; Gaps 1;

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QY 241 GLVPLPRADDSRLGVQVDTLHOVNOSSVOVLLFASVHAHAALFNYSISSRLSPKMWAS 300
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QY 421 VKPMQLLENMNLTHVGGPLRPFDSGQVNDMEYDLKLMWQGSVPRLHDVGRENGSLRT 480
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QY 481 ERLKIRMTSDNQKFEVSRCSROCOGOVRRVKGFSGCCYDVCCEAGSYRONPDDIACTF 540
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QY 541 CGODEMSPERSTRCFRRRSRFLAWGEPAYLILLILLISLALGLVLAALGLFVHHRDSPLVQ 600
DB 539 CGODEMSPERSTRCFRRRSRFLAWGEPAYLILLILLISLALGLVLAALGLFVHHRDSPLVQ 598
QY 601 ASGGPLACGLVCLVCLSVLLFPGQBPAPACLAQOPLSHPLTGCLSTLFLQAAEIFV 660
DB 599 ASGGPLACGLVCLVCLSVLLFPGQBPAPACLAQOPLSHPLTGCLSTLFLQAAEIFV 658
QY 661 ESELPLSWADRLSGCLRGFWAMLVYLLAMLEVALCTWYLVAFPEVVYTDHMLPTBALV 720
DB 659 ESELPLSWADRLSGCLRGFWAMLVYLLAMLEVALCTWYLVAFPEVVYTDHMLPTBALV 718
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DB 719 HCRTRSWVSFGIAHAATNATLAFCLGLTFVRSQPCYNRARGLTFAMLAIFYITWVSFVP 778

QY 781 LLANQVVLRPAVQWGAALLCIVGLIAAFHLPRCYLAKROPGNLTPEFFLGSGPDAGQ 840
DB 779 LLANQVVLRPAVQWGAALLCIVGLIAAFHLPRCYLAKROPGNLTPEFFLGSGPDAGQ 838
QY 841 NDGNTNGQKHE 852
DB 839 NDGNTNGQKHE 850

Search completed: May 11, 2004, 15:38:56
Job time : 44.3396 secs

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Qy	261	HOVNOSSVQVVLTFASVHAHAHLFNYSISSRLSPKVVASSEAWLTSDLVGLRPMQMGT	320
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Qy	321	VLGFLQRGQALHEFPQV-KTILALATDPAFCALGE-----	356
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Qy	357	REOGL-----EEDVQGRQRCOCCTLOVNSAGLNNHQFSPVAAVYSA	401
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Qy	402	QALHNTLOCNA-----SGCPAODEVKFMOULENNYNLTFFHV-GLPRLFDSGVDME	453
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Qy	454	YDLKLM--VMQGSVRLHDVGRFN-GSLRTERL-----KIRNHTDQKPVSRGCRQC	504
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Qy	505	EGQVVR-VKGFHSCCYDCVDCSAGSYRQNPDDICTFCGODEMSPSRSTRCFRRSRPLA	563
Db	547	PETRKGIIEGETTCCECVDCPDGEYSDETDASACCKCEBDYWSNENHTSCIPKQJEFIS	606
Qy	564	WGEPAVLLLLLLLSLALGLVLAALGLFVHHRDSPVQASGGPLA-----CFGLVCLGVCL-	619
Db	607	WTEPFGIALTLPAVIGIFLTSFVLGVTFRMNPVYKATNRBELSYLLFSLLC-----CFS	662
Qy	620	SVLLEPFGQSPARKLAQOPLSHULPTGCGSLTFLQAAELFV--ESELPLSMADRLSGCLR	677
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Qy	678	GPM-----AMLVLLAMLVEVALCTWYVLAAPPEVUTDWMHLMPTBAL-VHCRTRSWVSFG	711
Db	715	RKMGLNLOFLVLVFLCTFVQIYICVWLTAPRSSRN-HELEDEILFITCHEGSLMAG	773
Qy	732	LAHATNATLAFCLGTFLVRSQPCYNBARGLTFPAMLYFITWSPVFLLANVQVLR	791
Db	774	FLIGYCLLAAICFPFAFSRKLPEFNFAKFTTFMILFIWISIFIPAVASTYGGKVS	833
Qy	792	AVQMGALLCVGIIAALFHLPRCYLLMRQPLNTPRE	827
Db	834	AVEVIAIILAAISGLACIFPNKVYIILFPRSRNTIE	869

RESULT 2
US-08-485-588-7
Sequence 7, Application US/08485588
Patent No. 5688938
GENERAL INFORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hedbert
APPLICANT: Forrest H. Fallier
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESSES: 2

```

1 ADDRESS: Lyon & Lyon
2 STREET: First Interstate World Center
3 STREET: Suite 4700
4 STREET: 633 West Fifth Street
5 CITY: Los Angeles
6 STATE: California
7 COUNTRY: USA
8 ZIP: 90071
9
10 COMPUTER READABLE FORM:
11 MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
12 COMPUTER: IBM PC compatible
13 OPERATING SYSTEM: PC-DOS/MS-DOS
14 SOFTWARE: FASTSEQ
15
16 CURRENT APPLICATION DATA:
17 APPLICATION NUMBER: US/08/485,588
18 FILING DATE: 7 June, 1995
19
20 CLASSIFICATION: 435
21
22 PRIOR APPLICATION DATA:
23 PRIOR APPLICATION DATA: including application
24 PRIOR APPLICATION DATA: described below: 9
25 APPLICATION NUMBER: 08/353,784
26 FILING DATE: 9 December, 1994
27 APPLICATION NUMBER: PCT/US/94/12117
28 FILING DATE: 21 October, 1994
29 APPLICATION NUMBER: U.S. 08/292,827
30 FILING DATE: 23 August, 1994
31 APPLICATION NUMBER: U.S. 08/141,248
32 FILING DATE: 22 October, 1993
33 APPLICATION NUMBER: U.S. 08/009,389
34 FILING DATE: 23 February, 1993
35 APPLICATION NUMBER: U.S. 08/017,127
36 FILING DATE: 12 February, 1993
37 APPLICATION NUMBER: U.S. 07/934,161
38 FILING DATE: 21 August, 1992
39 APPLICATION NUMBER: U.S. 07/834,044
40 FILING DATE: 11 February, 1992
41 APPLICATION NUMBER: U.S. 07/749,451
42 FILING DATE: 23 August, 1991
43 ATTORNEY/AGENT INFORMATION:
44 NAME: Heber, Sheldon O.
45 REGISTRATION NUMBER: 38,179
46 REFERENCE/DOCKET NUMBER: 213/005
47
48 TELECOMMUNICATION INFORMATION:
49 TELEPHONE: (213) 489-1600
50 TELEFAX: (213) 955-0440
51
52 TELEX: 67-3510
53
54 INFORMATION FOR SEQ ID NO: 7:
55 SEQUENCE CHARACTERISTICS:
56 LENGTH: 1078 amino acids
57 TYPE: amino acid
58 TOPOLOGY: linear
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60 MOLECULE TYPE: protein
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62 US-08-485-588-7
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DB 299 WASSSLIAMPQYFHVVGITIGFALKAGQIPGFRFLKXVHPKSVHNGFAKFWETPNC 358
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DB 359 HLOEGAKGRLPYDTFLRGHEES--GDRFSNSSTAFRLCTGDENISSVETPYIDYTHLRI 416
QY 390 TFSVYAAYSVVAQALHNTLQ-----NASGCPADVPKPMOLLENNYULTF-HVGL 440
DB 417 STNVLIASVIAHALODITYTLPRGLFTNGS-CADIKVEAMQVILKHLNFTNNKE 475
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QY 608 CEGVLCLGLVCL-SVLLFPQSPARCLAQPLSHLPLTGCLSTLPLQAAEFV--ESEL 664
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QY 665 PLS-----WADRLSGCLRGPAMLVLLAMLEVALCTMYLVAPEPVTDNMLPTEAL 719
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RESULT 3

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US-08-484-565-7
; Sequence 7, Application US/08484565
; Patent No. 5763569
;
; GENERAL INFORMATION:
; APPLICANT: Edward M. Brown
; APPLICANT: Steven C. Hebert
; APPLICANT: James B. Garrett, Jr.
; TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Lyon & Lyon
; STREET: First Interstate World Center
; STREET: Suite 4700
; STREET: 633 West Fifth Street
; CITY: Los Angeles
; STATE: California
; COUNTRY: USA
; ZIP: 90071
;
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: FASTSEQ
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/484,565

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; FILING DATE: 7 June, 1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA: including application
; PRIOR APPLICATION DATA: described below: 9
; APPLICATION NUMBER: 08/353,784
; FILING DATE: 9 December, 1994
; APPLICATION NUMBER: PCT/US/94/12117
; FILING DATE: 21 October, 1994
; APPLICATION NUMBER: U.S. 08/292,827
; FILING DATE: 23 August, 1994
; APPLICATION NUMBER: U.S. 08/141,248
; FILING DATE: 22 October, 1993
; APPLICATION NUMBER: U.S. 06/009,389
; FILING DATE: 23 February, 1993
; APPLICATION NUMBER: U.S. 08/017,127
; FILING DATE: 12 February, 1993
; APPLICATION NUMBER: U.S. 07/934,161
; FILING DATE: 21 August, 1992
; APPLICATION NUMBER: U.S. 07/834,044
; FILING DATE: 11 February, 1992
; APPLICATION NUMBER: U.S. 07/749,451
; FILING DATE: 23 August, 1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Heber, Sheldon O.
; REGISTRATION NUMBER: 38,179
; REFERENCE/DOCKET NUMBER: 213/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 489-1600
; TELEFAX: (213) 955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO. 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1078 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
;
US-08-484-565-7

```

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Query Match 23.6%; Score 1068.5; DB 1; Length 1078;
Best Local Similarity 30.3%; Pred. No. 2.6e-93;
Matches 269; Conservative 156; Mismatches 366; Indels 97; Gaps 26;

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QY 12 WALL-----HPTGAPLCSQOLRMKGDYVGLGFLPGLG--EAEELGLSRTPSPVCTR 64
DB 8 WLLALTLWHTSAYGP---DORAQKKGDIIILGLFPIHGVAKDQDLKS--RPSEVECTR 62
QY 65 FSSNGLWALMKVAEEINNKSDLLPGLRLGYDLFDRCSEPVVMKPSIMPLA--KAGS 122
DB 63 YNFRGFRLOAVITFAIEBINSPALLPVLTLGRIPLFCNTVSVKALENTLSFVAQNKIDS 122
QY 123 RDIAYCNVTOYQOPRLAVVIGPHSSELAMVTGKPSFPLMQVSVGASMELLSARETPS 182
DB 123 LNLDFCNCHSEIIPSTIAVVGATGSGVSTAVANLLGLFYIIPQVYASSRLLSNNGQFSS 182
QY 183 FFRYVSPRVQVLTAAAEILQEFGNWVALASDDEYGRQGISIFSALAAAGICTAHESL 242
DB 183 FLRTIPNDEHQATMAADIIIEFRNMWVGITIAADDYGRGIEKFEAEAEERDICI DPFSEL 242
QY 243 VPLPRADSRGLKGVODVILHQNOSVOYVLLFPASVHAHALFNYSISSRLSPKVVASRA 302
DB 243 I-----SQSDEBEIQHVEVIONSTAKIVVSSGDEPLIKEIYRNITGKIWLASBA 298
QY 303 WLTSDLVNGLPGMAQMGTVLQFGAQLHEFPQYV-KTHLALATDPAFCALGERE--- 358
DB 299 WASSSLIAMPQYFHVVGITIGFALKAGQIPGFRFLKXVHPKSVHNGFAKFWETPNC 358
QY 359 -----QGLEEDVVGORCPQCD-----CITLQNVSA-----GLNHHQ- 389
DB 359 HLOEGAKGRLPYDTFLRGHEES--GDRFSNSSTAFRLCTGDENISSVETPYIDYTHLRI 416
QY 390 TFSVYAAYSVVAQALHNTLQ-----NASGCPADVPKPMOLLENNYULTF-HVGL 440

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Db 417 SYNVTYLAIVSAHALQDIYTCPLGRGLFTNGS--CADIKKVEAMQVLYKHLRHLNFTNNNGE 475
Qy 441 PLRFDSSGNVDMEDYDLKLM---VMQGSVPRLDHVGFRN--GSLRTERL-----KIRMTSD 491
Db 476 QVTFDECGDLVGNYSIINMHLSPEDGSIV-FKEVGYNVYAKGRLFINEKILMSGFS 534
Qy 492 NQKPVSRCSROCGQVRR-VKGFHSCCYDCVDCSAGSYRONPDIACFCQODEWSPER 550
Db 535 REVFPNSCSRDCLATRKQIIEGEPCTCCFCEVCEDPDEYSDETDASACNKCPCDDPMNEN 594
Qy 551 STRCFRRSRPLAWGEPAVLLLLLSLALGLVLAALGLFVHHRSPLVQASGGFLA--- 607
Db 595 HTSCIAKEIFLWTEPFGIALTLFAVIGIFLTAFLVGFIFKFRNPTPVKANNRSLSYLL 654
Qy 608 CFGVLGLGLVCL-SYLLFPQGPSPARCLAQOPLSHPLTGCTSTFLQAAEIFV--ESEL 664
Db 655 LFSLLC-----CFSSSLPFIQGEPODMTCRLRQAFISPLCLISCLVKTNRVLVPEAKI 710
Qy 665 PLS-----WADRLSGCLRGPMAMLVLLAMLVEVALCTWYLVAFPPREVTDHMLPTBAL 719
Db 711 PTFSRKMMWGLNLQ-----FLVPLCTFMQIVICVIMLYTADPSSYRNQLEDELIIF 762
Qy 720 VHCRRSRVSRFLAATNATLAFLCFLGTFVLVRSQPCYNRARGLTPMLAVFTWVSFV 779
Db 763 ITCHGSLMALGFLIGYTCCLAAICFPFAFKSRKLPENFNEMKFTTFSMILFFIWMISFI 822
Qy 760 PLLANVQVLRPAVOMGALLCGLAFLHPRCYLMLRQGLNTPPE 827
Db 823 PAYASTYKFSVAIVEVIALAASFGLLACIFPNKIIYIILFKSRNTIE 870

RESULT 4
US-08-480-751-7
Sequence 7, Application US/08480751

GENERAL INFORMATION:
APPLICANT: Edward F. Nemeth
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: Forrest H. Fuller
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCULON RECEPTOR-ACTIVE
TITLE OF INVENTION: MOLECULES
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/480,751
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993

APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/004
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-480-751-7
Query Match 23.6%; Score 1068.5; DB 2; Length 1078;
Best Local Similarity 30.3%; Pred. No. 2.6e-93;
Matches 269; Conservative 156; Mismatches 366; Indels 97; Gaps 26;
Qy 12 WALL-----HPGTGAPLCLSQLRMKGDYVLGGLFPLG---EABEAGLSRTRSPSPVCTR 64
Db 8 WVLLALTFMHTSAYGP---DQBAQKGDIIILGGLPPIHFGVAAKQDLKS--RPSVACIR 62
Qy 65 FSSNGGLMALAMKRAVEINNKSDLLPGLRLGDLPTTCGSEPPVYAMKPSLMFLA--KAGS 122
Db 63 YNFRGFPMLOAMIFAIEINSSPALLPNTLTYRIFPTCTVSKALATLTSFVAQNTIDS 122
Qy 123 RDLAAYCNYQYOPRVLAIVTGPSSSLAMTYGKFSPFLMPOVSYGASMETLSARETFPS 182
Db 123 LNLDFCNCSEHITSTIAVVGATSGVSTAVANILGLFYIPQVSYASSSRLLSKNOFKS 182
Qy 183 FFRTPSDRVQVLTAAAEILOEFGNNWVAALGSDDEYRGGLSIFSAALAAAGICIAHGL 242
Db 183 FLRTIPDEHQATMAADIIIEFRNNWGTIAADDDYRGPIEKREABEERDICTDPSL 242
Qy 243 VPLPRADSRGLKQVDVLHQVNOSSVOYVLLFASVHAHALFVNSISRLSPKVVASSEA 302
Db 243 I---SQYSDDEEIOHVVEVIONSTAKVIVFSSGPDLEPLIKEIVRNRITGKIMLASSEA 298
Qy 303 WLTSDLVWGLRGMNQMGTVLGFLOGAQLHEFPQVY-KTHLALATDPAFCALGERE--- 358
Db 299 WASSSLIAMPQYFHVVVGTTIGFALKAGQITPGFRFLKVKVHPRKSVHNGFAKEPWEETPNC 358
Qy 359 -----GLEBDVVGRCPOCD-----CITLQNVSA-----GLNHHQ- 389
Db 359 HLQGAKGRLPYDTFLRKGHESS--GDRFNSSTAFRLTCGDENISSVEMPYIDYTLRLI 416
Qy 390 TFSVYAAVSYVAQALNHTLOC-----NASGCPADPVKPMQLENNYTLTF-HVGL 440
Db 417 SYNVTYLAIVSAHALQDIYTCPLGRGLFTNGS--CADIKKVEAMQVLYKHLRHLNFTNNNGE 475
Qy 441 PLRFDSSGNVDMEDYDLKLM---VMQGSVPRLDHVGFRN--GSLRTERL-----KIRMTSD 491
Db 476 QVTFDECGDLVGNYSIINMHLSPEDGSIV-FKEVGYNVYAKGRLFINEKILMSGFS 534
Qy 492 NQKPVSRCSROCGQVRR-VKGFHSCCYDCVDCSAGSYRONPDIACFCQODEWSPER 550
Db 535 REVFPNSCSRDCLATRKQIIEGEPCTCCFCEVCEDPDEYSDETDASACNKCPCDDPMNEN 594
Qy 551 STRCFRRSRPLAWGEPAVLLLLLSLALGLVLAALGLFVHHRSPLVQASGGFLA--- 607
Db 595 HTSCIAKEIFLWTEPFGIALTLFAVIGIFLTAFLVGFIFKFRNPTPVKANNRSLSYLL 654

Db 823 PAYASTGKFSVAEVAIALAASGGLACIFPNKIYIILFKPSRNTIE 870

RESULT 6

US-08-353-784-7
Sequence 7, Application US/08353784
Patent No. 6011068

GENERAL INFORMATION:

APPLICANT: Edward F. Nemeth, Edward M.
APPLICANT: Brown, Steven C. Hebert,
APPLICANT: Bradford C. Van Wageningen, Manuel
APPLICANT: F. Balandrin, Forrest H. Fuller,
APPLICANT: Eric G. Delmar, and Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
NUMBER OF SEQUENCES: 20

CORRESPONDENCE ADDRESSES:

ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: FASTSEQ

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/353,784
FILING DATE: 9 December, 1994

CLASSIFICATION: 514

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 8

APPLICATION NUMBER: PCT/US/94/12117

FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827

FILING DATE: 23 August, 1994

APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993

APPLICATION NUMBER: U.S. 08/009,389

FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127

FILING DATE: 12 February, 1993

APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992

APPLICATION NUMBER: U.S. 07/834,044

FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451

FILING DATE: 23 August, 1991

ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179

REFERENCE/DOCKET NUMBER: 209/069

TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440

TELEX: 67-3510

INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:

LENGTH: 1078 amino acids

TYPE: amino acid
TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-353-784-7

Query Match 23.6%; Score 1068.5; DB 3; Length 1078;
Best Local Similarity 30.3%; Pred. No. 2.6e-93;
Matches 269; Conservative 156; Mismatches 366; Indels 97; Gaps 26;

Qy 12 WALL-----HPGTAPCLSQLRMKGDYVIGLEPLG---EAEFAGLRSTRPSSPVCTR 64
Db 8 WVLLALTMHTSAVGP---DQRAQKGDIIIGLGFPIHFVAAKQDILKS--RPFSEVCIR 62
Qy 65 FSSNGLLMALAKMAVEINNKSDLLPGLRLGDLPTTCSEPPVAMKPSLMFLA--KAGS 122
Db 63 YNFRGFRLQAMFAISEINSSPALPLNLTLGYRIFDTCNTVSKALEATVSPVAQNIDIS 122
Qy 123 RDLAAYCNQYQYPRVAVAVIGPHSSELMATYGFSSFFLMPQVSYGASMEILSAREFPS 182
Db 123 LNLDFPCNCEHPISTIAVAVGATGSSVSTAVANLGLFYIPQVSYASSSSLLSKNQPKS 182
Qy 183 FFRTPSDRVQVLTAAELDLQFGNNVVAAGSDDEYGRQGLSTFSAALAAAGICIAHGL 242
Db 183 FLRTIPNDEQATAMADIIIEYFRNWNVGTIAADDYGRPGIEKREBAEERDIDIPSEL 242
Qy 243 VPLPRADDSRLGKQVDTLHOVNOSSVQVLLFPASVHAALHFNYSISRLSPKYMVSEA 302
Db 243 I---SQYSDSEELQHVAVIQNSTAKVIVFSSGPDLEPLIKEIVARNITGKIWLSEA 298
Qy 303 WLTSDLVWGLPGMAQMGTVLGFLOGAQLHEFPQYV-KTHLATATDPAFCALGERE--- 358
Db 299 WASSSLAMQYFRVAVGTTIGFALKAQOIPGFRFLKQVPRKS VNHGFAKEFEETPNC 358
Qy 359 -----QGLEDDVVGORCPQCD-----CITLQNVSA-----GLNHHQ- 389
Db 359 HLOGAKGPLPVDFTFLRGHEES--GDRFNSSTAFRLCTGDEISSVETPYIDYTLRI 416
Qy 390 TFSYVAAVSVAAQNLNHTLOC-----NASGCPADDPKPMOLENMNLTFF-HVGL 440
Db 417 SYNVYLAIVYSIAHALQDIYTCLPGRGLFTNGS--CADIKKVAQVNLGHLNHTNMGE 475
Qy 441 PLRFDSSGNVDMEXDLTM---VWQGSVPLRHDVGRFN--GSLRTERL-----XIRWTS 491
Db 476 QVTFDECGDLVGNYSIINMHLSPEDGSIV-FKEVGYNVVAKKGERLFIIBEKILWGSFS 534
Qy 492 NQKPVSCSCQOQEGQVVR-VKGFHSCCYCDVCEAGSYQNPDDIACCTCGODESPER 550
Db 535 REVFPFNSCSDCLAGTKKGIIEGEPCTCEPCVECDPDEYDSDTASACNCPDPFWSNEN 594
Qy 551 STRCFRRRSRFLMGEBAVLLLLLSLALGVLAALGLFVHHRDSPLVOASGGPLA--- 607
Db 595 HTSCIAIEPLSTWTEPFGIALTLFAVLGIFLTAFLVGLVTKRNTIYATNRELSYLL 654
Qy 608 CFGLVCLGLVCL-SVLLPQGPSPARCTAQQPSHLPTCLSTLFLQAEIFV--ESEL 664
Db 655 LFSLLC-----CFSSSLFFIEBPQDMTCRLQRPAGISFVLCIGCILVKNRVLVFEAKI 710
Qy 665 PLS-----WADRUSGCLRGPMWLVLVLLAMLEVALCTWYLVAFPPREVTDHMLPTEAL 719
Db 711 PTFEHRKMGMLNQ-----FLVPLCTFMQVICVILWYAPRPSYRNQELEDEIIF 762
Qy 720 VHCRTRWVSFGLAHATNATLAPLCFLGTFLVNSQPCVYRARGDTFAMLAFTWVSFV 779
Db 763 ITCHEGSLMALGFLIGYTCLLAALCFEPFAKSRKLPENFNEAKRITFSMLIFLWISFI 822
Qy 780 PLANOVVLRPAVOMGALLCVLGILAFHLPRCYLAKRQPLNTPPE 827
Db 823 PAYASTGKFSVAEVAIALAASGGLACIFPNKIYIILFKPSRNTIE 870

RESULT 7

US-08-484-719B-7
Sequence 7, Application US/08484719B
Patent No. 6031003

GENERAL INFORMATION:

APPLICANT: Edward F. Nemeth, Edward M.
APPLICANT: Brown, Steven C. Hebert,
APPLICANT: Bradford C. Van Wageningen,
APPLICANT: Manuel F. Balandrin,
APPLICANT: Forrest H. Fuller, Eric G.
APPLICANT: Delmar, Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE

```

; TITLE OF INVENTION: MOLECULES
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Lyon & Lyon
; STREET: First Interstate World Center
; STREET: Suite 4700
; STREET: 633 West Fifth Street
; CITY: Los Angeles
; STATE: California
; COUNTRY: USA
; ZIP: 90071
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: MS Word
; SOFTWARE: FastSeq for Windows Version 3.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/484,719B
; FILING DATE: 7 June, 1995
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/353,784
; FILING DATE: 9 December, 1994
; APPLICATION NUMBER: PCT/US/94/12117
; FILING DATE: 21 October, 1994
; APPLICATION NUMBER: U.S. 08/292,827
; FILING DATE: 23 August, 1994
; APPLICATION NUMBER: U.S. 08/141,248
; FILING DATE: 22 October, 1993
; APPLICATION NUMBER: U.S. 08/009,389
; FILING DATE: 23 February, 1993
; APPLICATION NUMBER: U.S. 08/017,127
; FILING DATE: 12 February, 1993
; APPLICATION NUMBER: U.S. 07/934,161
; FILING DATE: 21 August, 1992
; APPLICATION NUMBER: U.S. 07/834,044
; FILING DATE: 11 February, 1992
; APPLICATION NUMBER: U.S. 07/749,451
; FILING DATE: 23 August, 1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Douglas C. Murdock
; REGISTRATION NUMBER: 37,549
; REFERENCE/DOCKET NUMBER: 213/007
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 489-1600
; TELEFAX: (213) 955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1078 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-484-719B-7.

```

Query Match 23.6%; Score 1068.5; DB 3; Length 1078;
 Best Local Similarity 30.3%; Pred. No. 2.6e-93;
 Matches 269; Conservative 156; Mismatches 366; Indels 97; Gaps 26;

```

QY 12 WALL-----HPTGALCLSCQDRMGDYLGLPPLG---EAEAGLGRTRTPSSPVCTR 64
DB 8 WLLALLTHTTAYGP---DORAKKGDDIILGLFPIHGVAKDDOLKS--RPESVEICR 62
QY 65 FSSNGLLWALAKMAVEEINNKSDLLPGILGGLDFTCSBPVAMKSLMFLA--KAGS 122
DB 63 YNFRGFRLQAMIFAIIEINSSPALLPNTLIGYRIFDTCNTVSKALEATLSFVAQNKIDS 122
QY 123 RDIAYCNYTQYQRPVLAVTGHSSSLAMVTKGKFSFPLMPQVSYGASMEILISAETPPS 182
DB 123 LNLDFCNCSEHIFPTIAVVGATGVSVAVALNLGLFPIPOVSYAASSSRLLSNKQFYS 182
QY 183 FFRVPSDRVQLTAALLLQERFGMMVVALGSDDEYGRGSLISFSLAAARGICIAHGL 242

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DB 183 FLRTIPNDHQATAMADIIIEYFRMNVGTIAADDDYGRPGIEKFEAEAEEDIDICIDFSEL 242
QY 243 VPLPRADSRKQKQVDVHQNQSSVQVLLPASHAALFNYSISSRLSPKVVASSEA 302
DB 243 I---SQYDDEEIOHVEVIQNSTAKVIIVFSSGPDLEPLIKETIRNITKMLASSEA 298
QY 303 WLTSDLVNGLPQMAQMGTVLGLQGAQLHFPPQV--KTHLALATDPAFSCALGERE--- 358
DB 299 MASSSLIMPQYFHVVGSTIGFALKAGQIPGFRERLKVHVRKSHNCFALFEMETENC 358
QY 359 -----QGLEBDVVGQRCPCD-----CITLQNVSA-----GLNHHQ- 389
DB 359 HLEGAKGKPLPVDFELRGHEES--GDRFSNSTAFRLPCTGDEINISVETPIDYTHLRI 416
QY 390 TFSVYAAIYSAQALHNTLQC-----NASGCAQDPVYPMOLEMMYLTFF-HVGL 440
DB 417 SYNVLAVYSIAHLAQDITYCLPGRLFTNGS-CADIKVAMQVLRKALHNFNNNGE 475
QY 441 PLRFDSSGNVDMEXDLKLM---VMQGSVRLHDVGRFN--GSLRTERL-----KIRWHTSD 491
DB 476 QYTFDECGDLVGNISITMHLSPEDGSIV-FKEVGYYVYAKKGERLITNEKILMSGFS 534
QY 492 NQKPVSRCSROCEQOYRR-VKGFHSCCYDCVDCBAGSYRONPDDIACFPQODEWSPER 550
DB 535 REVPSNCSRDCLAGTRGIIIEGPTCCFECVCEGDSYSETDASACNKKCPDDFWSMEN 594
QY 551 STRCFRRSRFLANGEPVALLLLLSLALGLVLAALGFVHHRSDPLVQASGGLA--- 607
DB 595 HTSICAKIEIEFLSWTEPPGIALTLFAVLGIFLTAFLVGLVFTRNTPIVKAITNELSYLL 654
QY 608 CEGVLCGLVCL-SVLPFRGQSPARCLAQPLSLPLTGLSTLFLQAALIFV--ESEL 664
DB 655 LPSLCL---CPSSLFPTGEPQDWTCKLRPARGISVLTLSCLVLTNRLLVFEAKI 710
QY 665 PLS-----MADRLSGCLRGPMAMLVLLAMVEVALCTWYLVAPPEVYVTDHMLPTBAL 719
DB 711 PTFHRKMWGMLNQ-----FLVFLCTMQIVICWLYTAPSSYRNGELEDLEIF 762
QY 720 VHCRTSRVNSFGLAHATNATLAFLCFLGTFLVRSOPGCTYRARGLTFLMAYFTVWSPV 779
DB 763 ITCHEGSLMAGFLIGYTCLLAALCFEPFAFSRKLPEFNFAKFTFPMLLFFIWMISFI 822
QY 780 PLIANVQVVLPAVOMGALLCVGIIAFLPLPCYLLMRQGLNTPPE 827
DB 823 PAYASTYKFSVAVEVAILLAASFGLACIFPNKIYIILFRPSRNTIE 870

```

RESULT 8
 US-08-484-159-7
 ; Sequence 7, Application US/08484159
 ; Patent No. 6313146
 ; GENERAL INFORMATION:
 ; APPLICANT: Bradford C. Van Wageningen
 ; APPLICANT: Manuel F. Baladrin
 ; APPLICANT: Eric G. Del Mar
 ; APPLICANT: Edward F. Nemeth
 ; TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
 ; MOLECULE TYPE: MOLECULES
 ; NUMBER OF SEQUENCES: 20
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: Lyon & Lyon
 ; STREET: First Interstate World Center
 ; STREET: Suite 4700
 ; STREET: 633 West Fifth Street
 ; CITY: Los Angeles
 ; STATE: California
 ; COUNTRY: USA
 ; ZIP: 90071
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS

```
SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,159
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 9
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Hebert, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 214/101
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 1078 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-484-159-7

Query Match      23.6%  Score 1068.5; DB 4; Length 1078;
Best Local Similarity 30.3%  Pred. No. 2.6e-93;
Matches 269; Conservative 156; Mismatches 366; Indels 97; Gaps 26;

12 WALL---HPTGALCLISQULRMGDYVLGSLFPLG---EAEZGLASRTSPSPVCTR 64
8 WVLALTYHTSAVGP---DORAQKGDIIIGGLPIHFGVAKDODLKS--RPESVEGIR 62
65 PSSNGLMALMKMAVEREINNKSDLLPGLRIGYDFPTCSEBVVMKPSLMPFLA--KAGS 122
63 YNFRGFRWLQAVIPIEINSSPALLPMTLIGYRIPTCNVYSKALETLSTVAONKIDS 122
123 RDIAVVCNYTOYQPRVLAVIGPSSSELAMVTGKFSFLMPQVSYGASMEELISARETPS 182
123 LNLDFCNCSEHIEPTIAVVGATGVSFAVANLLGLFIYIPQVSVASSRLLSNNGQKFS 182
183 FFRYVPSRVOLTAABELLOEFGMNWVAALGSDDEYGRQGLSIFALAAAGICIAHGL 242
183 FLRTIPNDEHQTAADIIYFRMNWVGIIAADDYGRPGIEKFEFEAEERDICTDFSEL 242
243 VPLPRADSRIGKVDVLIHQVNOSSVOVLLFASVHAHAALFNYSISRLSPKVVAASEA 302
243 I---SQYDEBEIGHVEVIONSTAKVIVFSSGPDLEPLKEIVRNNTIGKMLASEA 298
303 WLTSLDVLWGLPGMAQMTVLGFLQGAOLHEFPQV-XTHLALATDPAFCSALGERE--- 358
299 MASSSLIAMPQYFNHVVGIGTIFALAAAGIIPGFRRELKVKVHPKSKVHNGFAKFWMETENC 358
359 -----QGLEEDVVCORCPQCD-----CITLQNVSA-----GINHQ- 389
359 HLOGAKGKPLPVDTFLRGHEES--GDRFNSNSTAFRPLCTGDENISSVETPYIDYTHIRI 416
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390 TFSYAAVSYVAQALNHTLQC-----NASGCPADPVKPMQLENNVLTFF-HVGC 440
417 SYNYLVAVYSLAHALDQIYTCPLRGFLFTNGS-CADIKYBAWQVLIHRLHNTNNMGE 475
441 PLRTDSSGNVMEYDLKM--VMQSVPRLDHYGRN-GSLRTERL-----KIRHNTSD 491
476 QVTEDECGDVLGNYSIINMHLSPEDGSIV-FKEVGYNVYAKGGERLFINBEKILMGFS 534
492 NOKFVSRCSQCQCGOYVR-VKGFHSCCYOCVDEAGSYRONPDDICTFCGODEWSPER 550
535 REVPSNCSRDCLAGTKGIIIEGPTCCFECVBCPDDEYSDETDASCKNCCPDPMSENEN 594
551 STRCFRRSRPFLANGEPVALLLLLSLALGLVLAALGLFVHNHSDPLVQASGGPLA--- 607
595 HTSCIAKEIEFLSWTEPFGIALTLFANVLGIFLTFVGLVGFPIKPNNTIIVATNRELSYL 654
608 CFGVLVCLGLVCL-SVLLPQGPSPARCLAOQPLSLPLTGLSLTLFLQAAEIFV--ESEL 664
655 LFSILC-----CFSSSLPFIIGBPQDMTCRLROPAGISFVLICISILVKTNRVLLVPSAKI 710
665 PLS-----MADRISGCLRGFWAMTLVLLAMLVVALCTWYLVAFRPREVTDMHMLPTREAL 719
711 PTFSHRRKFWGLNLD-----FLVFLCTFMQVIVICVIMLYTAPBSYRNQELDEDTIF 762
720 VHCRTSRWVSFGLAHATNATLAFPLCFLGTFLVRSQPCYNRARGLTPEMLAYFITWVSFV 779
763 ITCHEGSLMALGFIIGTCTLAAICFPFAFYSRKLPENFDEAKITFSMLIFLWISFI 822
823 PAVASTGKEVSAVEVIAIILASFGILACIFFNKIYIILKPSRNTIE 870

RESULT 9
US-08-485-588-5
Sequence 5, Application US/08485588
Patent No. 5688938
GENERAL INFORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: Forrest H. Fuller
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,588
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 9
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
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APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 213/005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ. ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1085 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-485-588-5

Query Match 23.4%; Score 1060; DB 1; Length 1085;
Best Local Similarity 30.0%; Pred. No. 1.8e-92;
Matches 269; Conservative 155; Mismatches 362; Indels 112; Gaps 26;

6 VGLSLMLHRTGAPLCLSLQGLMKGDYVIGLPL--GEA-EEAGLRSTRSSPVC 62
10 LIAFTWCTSAVGP-----DQRAQKGIILGGPLHFGVAVXDQDKS--REPSVVC 61
63 TRFSSNGILMALAMKAAVEINNKSDLLPGLRIGYDLFTGCEPVAAMKPSIMFLA--KA 120
62 IRYNRRGRWLOAMFAIEEINSSPALLPNNMLGRIPTCTVSKALEATLSFAQNKI 121
121 GSRDIAVCNTYQYOPRVLAIVIGPHSSSELAIVTGKFFSEFLMPQVSYGASMELLARETF 180
122 DSLINDEFONCESEHPISTIAVAVGATGSGISTAVANLLGFIYIPQVSYASSSRLSNKQF 181
181 PSFRTVSDRYQLPAAAELOEPGMNVAALGSDDEVROGLSTFSALAAARGICIAHE 240
182 KSFLLTIPDEHQATAMADILEYFRMNWGTIAADDDYGRPGIEKFRREAEERDICIPTS 241
241 GLVPLPRADDSRLGKQVIVLHQNOSVQVLLPFAVHAHALFNYSISSRLSPKVVAS 300
242 ELI-----SOYDEBEKIQVVEIYQNSTAKVIYVSSGPLELEIYVRNITGRILWAS 297
301 EAMLTSDLVMLPGNAQGTIVLGLRGAQLHEPQYV-KTHLALATDPAFCAL----- 354
298 EAMASSSLIAMPDEYFHVVGITIGFGKAGQIPGFRFLOKVAPRKSVMHNGFAKEFEWETF 357
355 -----GEREGQL-----BEDVVGRCPCQDCITLON 380
380 NCHLOEAGKPLPVDFTLRGHEEGGARLSNSPFAAPRLCTGSEINSVETPMDYTHLR 416
381 VSAGLHHQTFESVAAYVSAQALHNTLOC-----NASGCPADPVKPKQLLENMNY 432
417 -----ISYVYLAIVSIAHALODIYTCIRGRGLFTNGS-CAIKKYEAHQVLAHLRH 467
433 LTFHVG-GLPLRFDSGGVNDMEYDLKW---VMQGSVPLRLHDVGRFN-GSLRTERL--- 483
468 LNFSTNMGEQVTFEDCGDLAGNYSIINWHLSPEDSIV-FKEVGYVNAVAKGGERLFIN 526
484 -KIRWHTSDNOKVRCSCROGEGQVRR-VKGFHSCVDVCVCEKSGYRQNDDDIACIFC 541
527 EXILWSGSRREVPSFNSCSRDCLAGTRKGIIEBEPFCCEFCVCPGDEESDETASACDKC 586
542 GODEWSPERSTRCFRRSRFLAMGEPVAVLLILLLSLALGLVLAALGLFVHHRDSPLYQA 601

DB 587 PDDFWSNENHTSCIAKEIEPLSWTEPFGIALTLFVAVLGFPLFAVFLGVFIKFRNTPYVA 646
QY 602 SGGPLA---CGVLVLGLVCL-SVLLFPGQSPAPACLAQOPLSHPLRGCLSTLFLQAAE 657
DB 647 TNRRLSYLLLSFLCL-----CFSSSLFPIGEPDMWTCRQRPAGFSGFVLCISCIILVKNR 702
QY 658 IFV--ESELPS-----WADRLSGCLRPMMLVYLLMLVVALCTWVLAFFPEVVD 710
DB 703 VILVEBAKIPTSFHRKMGMLNQ-----FLVFLCTFMQIVICAIWLTAPSSSYRN 754
QY 711 WHMLPTEAL-VHCRTRSVSFGLAHATNATLAFCLGTLFVRSQPGCVNRRARGLTFAML 769
DB 755 -HELEDEILFIICHGSLMALGFLIGYCLLAALCFEFAFSRKLPENNNAKFTFSML 813
QY 770 AYFITVSGFVPLANVOVVLPAVOMGALLLCVLGILAAFHLPRLCYLLMROGLTAPTE 827
DB 814 IFFIVWISFIPAVASTYGVSAVEVIAIILAAASFLLACIFENKVIILFKPSRNTIE 871

RESULT 10
US-08-484-565-5
Sequence 5, Application US/08484565
Patent No. 5763569
GENERAL INFORMATION:
APPLICANT: Edward M. Brown
APPLICANT: Steven C. Hebert
APPLICANT: James E. Garrett, Jr.
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: First Interstate World Center
STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,565
FILING DATE: 7 June, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 9
APPLICATION NUMBER: 08/353,784
FILING DATE: 9 December, 1994
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
FILING DATE: 12 February, 1993
APPLICATION NUMBER: U.S. 07/934,161
FILING DATE: 21 August, 1992
APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Heber, Sheldon O.
REGISTRATION NUMBER: 38,179

```

; REFERENCE/DOCKET NUMBER: 213/006
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 489-1600
; TELEFAX: (213) 955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1085 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-480-751-5

Query Match      23.4%; Score 1060; DB 1; Length 1085;
Best Local Similarity 30.0%; Pred. No. 1,8e-92;
Matches 269; Conservative 155; Mismatches 362; Indels 112; Gaps 26;

QY 6 VLGLSLMALHPTGAPRLCSQQLRMKGDYVGLFPL--GEA--EAGLRSTRPSPVQC 62
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 10 LLAFTWCTSAVGP-----DQRAQKKGDIIIGLFPFHGVAVKDDLKS--RRESVEC 61
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 63 TRPSSNGLLMALAMMAVEIRINKSDLLPGLRLGIDLPDTGSEPVVAMKPSLMLPLA--KA 120
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 62 IRYNFRGRFWLQAMFAIEEINSSPALLPNNMTLGRIIDTCNTVSKALEATLSFAAQYKI 121
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 121 GSRDIAACNTYQOPRVLAIVGPHSSSELAMVTKGFSPFLMPQVSYGASMELSARETF 180
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 122 DSLNIDECNCSEHPTITAVGARGSGISTAVANLGLFYIPQVSYASSRLSNKQF 181
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 181 PSFRTVPSDEVOLTAABELLOEFGMNVAALGSDDEXYRGQGLSFSLAARGICIAHE 240
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 182 KSPFLRTIYNDHQATAMADIIEYFRMNWVGITAADDDYGRGIEKFEAREEERDIDFS 241
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 241 GLVPLPRADBRGLGVQVNLQVNSSVQVNLFPASVAALAPNYSISSLSPRVWAS 300
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 242 ELI---SQYSDDEKIQOVVEVIONSTAKVIVFSSGPDLEPLKEIYRNITGIMLAS 297
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 301 EAMLTSDLVMLPGMAQMGTVLGFLOGAQLHEFPQVY--KTHLALATDPAFCSAL----- 354
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 298 EAMASSSLIAMPYRHVVGITIGPLKAGQIRGPRFEPLOXNHRKSVANGAKKEWETTF 357
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 355 -----GERREGQL-----BEDVVGRCPCQDCDTYLQN 380
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 358 NCHLOEGAKGPLVVDTPLRGHEGEGARLSNPAPRPLCTGEENISSVETPYMDYTHLR 416
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 381 VSAGLNHTQTSYVAAYVAQAALHNTQC-----NAGGCRAPQDVKKWOLLENNYN 432
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 417 -----ISYNYLAIVSIAHALODIYTCIPRGGLFTNGS--CADIKKVEAMQVULKLRH 467
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 433 LTFPHVG--GLPLRPDSSGVNDEMYDYLKLW---VMQGSVRLHDVGRFN--GSLRTERL----- 483
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 468 LNFISNMGEQYLPFECBDLADGNYSIINMHLSPEDSIV--FKEVGVYNYAKKGEHLFIND 526
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 484 -KIMHTSDNOKPVSRGRCQCOEGOVRR--VKGPHSCCYDCVDCESGRONPDIACFFC 541
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 527 EKILMSGSRVREPVSNCGRCDLAGTRKGIIEGEPTCECEVCPEGSEYSDETDASACDKC 586
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 542 GODEMSPRSRTRCPRRRSRFLAMGRPAVLULLLLLSLGLVLAALGFVHHRSPLVQA 601
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 587 PDDFWSNHNHTSCIAKEIEFLSWTEPGIATLFLPAVLGIFLPAVLGVFIERNTPVYKA 646
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 602 SGGPGLA---CFGLVCLGLVCL--SVLLPFGOPSPARCLAQOPLSHLPLGLCSTLFLQAAE 657
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 647 TNRELSYLLFSLIC-----CFSSSLFFGEPQDWCRLRGARFAGSFLVCLISCLIVKNR 702
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 658 IFV--ESSELPLS---WADRLSGCLRPAMLVLLAMLVEVALCTWYLVAPPEVVD 710
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 703 VLVLEAKIPTSPFRHKWGLNMQ-----FLVFLCTFQIYCAIWLNTAPSSRYRN 754
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
QY 711 WHMLPTEAL--VHCRRSVSVFGLAHATNATLAFLCFLGTFPLVRSGPGCVNARGLTFAML 769
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 755 -HELEDEITFTCHGSLMALGFLIGYCLLAALICFFPAFASRKLPENFNAKKTFTFSL 813

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QY 770 APTFTVSVFVLLANVQVILRPVAVMGALLCVLIGILAAFLPRCYLLMRQGLNTPB 827
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:
DB 814 IFFIWTISFIPAVASTGKFSVAVEVIALAASFGLACIFPNKNVYIILFKPSRNTIE 871
   |||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:|||||:

RESULT 11
US-08-480-751-5
; Sequence 5, Application US/08480751
; Patent No. 5856684
; GENERAL INFORMATION:
; APPLICANT: Edward F. Nemeth
; APPLICANT: Edward M. Brown
; APPLICANT: Steven C. Hebert
; APPLICANT: Forrest H. Fuller
; APPLICANT: James E. Garrett, Jr.
; TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
; NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Lyon & Lyon
; STREET: First Interstate World Center
; STREET: Suite 4700
; STREET: 633 West Fifth Street
; CITY: Los Angeles
; STATE: California
; COUNTRY: USA
; ZIP: 90071
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: FASTSEQ
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/480,751
; FILING DATE: 7 June, 1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; PRIOR APPLICATION DATA: including application
; PRIOR APPLICATION DATA: described below: 9
; APPLICATION NUMBER: 08/353,784
; FILING DATE: 9 December, 1994
; APPLICATION NUMBER: PCT/US/94/12117
; FILING DATE: 21 October, 1994
; APPLICATION NUMBER: U.S. 08/292,827
; FILING DATE: 23 August, 1994
; APPLICATION NUMBER: U.S. 08/141,248
; FILING DATE: 22 October, 1993
; APPLICATION NUMBER: U.S. 08/009,389
; FILING DATE: 23 February, 1993
; APPLICATION NUMBER: U.S. 08/017,127
; FILING DATE: 12 February, 1993
; APPLICATION NUMBER: U.S. 07/934,161
; FILING DATE: 21 August, 1992
; APPLICATION NUMBER: U.S. 07/834,044
; FILING DATE: 11 February, 1992
; APPLICATION NUMBER: U.S. 07/749,451
; FILING DATE: 23 August, 1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Heber, Sheldon O.
; REGISTRATION NUMBER: 38,179
; REFERENCE/DOCKET NUMBER: 213/004
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (213) 489-1600
; TELEFAX: (213) 955-0440
; TELEX: 67-3510
; INFORMATION FOR SEQ ID NO: 5:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1085 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-480-751-5

```


Query Match 23.4%; Score 1060; DB 2; Length 1085;
 Best Local Similarity 30.0%; Pred. No. 1.8e-92;
 Matches 269; Conservative 155; Mismatches 362; Indels 112; Gaps 26;

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QY 6 VIGLSMALHPGTGAPLCLSQLRMKGDYVIGLFLP--GEA--EAGLRSTPRSSVYC 62
DB 10 LIAFSTWCTSAVGP-----DORAKKGDIIIGLGFPIHFGVAVVDQDLKS--RESVEVC 61
QY 63 TRFSSNGLLMALAMKMAVEINNKSDLLPGLRLGYDLPDTCSEBPVAMKPSLMFLA--KA 120
DB 62 IRVNRGRFRLQAMIFALEINSSPALPNNLTGLRIPTDTCNTVSKALBATLSFPAQNKI 121
QY 121 GSRDIAAYCNYTOYOPRVAVIGPHSSSELAWTKGFSSFFLMPQVSYGASMEILLSARETF 180
DB 122 DSLNIDFCNCSEHLPSTIAVVGATGSGSTAVANLGLFYIPQVSYASSSRLLSKNQF 181
QY 181 PSFFRTVSDRVQLTAAELLOEFGNNVVALGSDDEYRGQSLSFSAALAAARGICIAHE 240
DB 182 KSFLRTIPVDEHQATAMADIIIEFRMNVGTIAADDYGRPGIEKFREREABERDICIIDS 241
QY 241 GLVPLPRADDSRLGKVODVLHQVNOSSVQVVLFPASVAAHALLFNYSISSRLSPKWTAS 300
DB 242 ELL-----SOYSDEEKIQOVVEVIONSTAKVIVFSSGPDLEPLKEIVARNITGRILWLS 297
QY 301 EAMLTSDVLMGLPGMAOMGTVLGFLQGAQLHEFPQYV-KTHLALATDPAFCAL----- 354
DB 298 EAMASSSLJAMPEYHVVGGTIGFGLKAGQIGFREFLOKHPKRSYVNHGFAKEWERTF 357
QY 355 -----GEREQL-----BEDVQGRCPQCCDITLON 380
DB 358 NCHLOEAKGKPLPVDTLRGHEEGARLSNSPTAFRPLCTGEMNISVETPYMDYTHLR- 416
QY 381 VSAGLNHHQTESVYAAVSVAAQLNHLNQLQ-----NASGCPADDPKPKQMLLENMYN 432
DB 417 -----ISVNYAVYSIAHALDITYTCLRGKFLTNCS--CADIKKEAQVKKLHLH 467
QY 433 LTFHVG-GLPLRFDSGNDMEYDKLW---WQGSVPLRHVQGFN--GSLTEPL----- 483
DB 468 LNFNTMGBQVTFDECGDLAGNYSIINMHLSEDSISV-FKEVGYVYVAKKGERLFIN 526
QY 484 -KIRNHTSDNQKPVSRCSRQCEGOVRR-VKGFHSCCYDCVDCGASGRQNDPIACTFC 541
DB 527 EKILMSGFSREVPFNCSDCLAGTRKGIIEGEPICCEFCVCPGDEYDETDASACDCK 586
QY 542 GODEMSPERSPTCFRRSRFTLWGPBAYVLLLLLSLALGVIALGLFVHHRDSPVQA 601
DB 567 PDDFWSNENHTSCIAKEIEFLSWTEPPGIALTLFVGLIFLTAFLVGVIKFRNTPVYA 646
QY 602 SGGPLA---CFGLVCLGLVCL-SVLLFPQPSPARCLAQOPLSHLPLTGCLSTLFLQAAE 657
DB 647 TNRELSTYLLLFELLC-----CFSSSLFFIGEPQDWTCLRQPAFGISFVLCISCIIVKTR 702
QY 658 IFV--ESELPLS-----WADRLSGCLRGFWMLVVLAMLVVALCTWYLVAPPEVYTD 710
DB 703 VLLVFEAKIPTSFHKKMGLANIQ-----FLAVELCTFMQIVICAIMLNTAPSSRYN 754
QY 711 WHMLPTEAL-VHCRTRSWSPGLAHATNATLAFLEPLGLFVLRSQPGCINRARGILTEPML 769
DB 755 -HELEDEIIFITCHEGSLMALGFLIGYTCLLAALCFEFAFKSRKUPENNENKFTTFSL 813
QY 770 AVFITWGFVPLLANVQVVLRAVOMGALLLCVLGILAAFLHPRCVLMLRQGLTPE 827
DB 814 IFFIWMISFIPAYASTYKGFVSAVEYIALAASFGLLACIFPNKYIIILFKSRNTIE 871

```

Query Match 23.4%; Score 1060; DB 2; Length 1085;
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 Matches 269; Conservative 155; Mismatches 362; Indels 112; Gaps 26;

```

QY 6 VIGLSMALHPGTGAPLCLSQLRMKGDYVIGLFLP--GEA--EAGLRSTPRSSVYC 62
DB 10 LIAFSTWCTSAVGP-----DORAKKGDIIIGLGFPIHFGVAVVDQDLKS--RESVEVC 61
QY 63 TRFSSNGLLMALAMKMAVEINNKSDLLPGLRLGYDLPDTCSEBPVAMKPSLMFLA--KA 120
DB 62 IRVNRGRFRLQAMIFALEINSSPALPNNLTGLRIPTDTCNTVSKALBATLSFPAQNKI 121
QY 121 GSRDIAAYCNYTOYOPRVAVIGPHSSSELAWTKGFSSFFLMPQVSYGASMEILLSARETF 180
DB 122 DSLNIDFCNCSEHLPSTIAVVGATGSGSTAVANLGLFYIPQVSYASSSRLLSKNQF 181

```

TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
 TITLE OF INVENTION: MOLECULES
 NUMBER OF SEQUENCES: 20
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Lyon & Lyon
 STREET: First Interstate World Center
 STREET: Suite 4700
 STREET: 633 West Fifth Street
 CITY: Los Angeles
 STATE: California
 COUNTRY: USA
 ZIP: 90071
 COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: FASTSEQ
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/943,986
 FILING DATE: 03-OCT-1997
 CLASSIFICATION: 530
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: 08/484,565
 FILING DATE: 7-June-1995
 APPLICATION NUMBER: 08/353,784
 FILING DATE: 9 December, 1994
 APPLICATION NUMBER: PCT/US/94/12117
 FILING DATE: 21 October, 1994
 APPLICATION NUMBER: U.S. 08/292,827
 FILING DATE: 23 August, 1994
 APPLICATION NUMBER: U.S. 08/141,248
 FILING DATE: 22 October, 1993
 APPLICATION NUMBER: U.S. 08/009,389
 FILING DATE: 23 February, 1993
 APPLICATION NUMBER: U.S. 08/017,127
 FILING DATE: 12 February, 1993
 APPLICATION NUMBER: U.S. 07/934,161
 FILING DATE: 21 August, 1992
 APPLICATION NUMBER: U.S. 07/834,044
 FILING DATE: 11 February, 1992
 APPLICATION NUMBER: U.S. 07/749,451
 FILING DATE: 23 August, 1991
 ATTORNEY/AGENT INFORMATION:
 NAME: Hebert, Sheldon O.
 REGISTRATION NUMBER: 38,179
 REFERENCE/DOCKET NUMBER: 213/006
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (213) 489-1600
 TELEFAX: (213) 955-0440
 TELEX: 67-3510
 INFORMATION FOR SEQ ID NO: 5:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 1085 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-943-986-5

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QY 181 PSFRTVPSDRVQVLTAAABELOEFQGMNVAALGSDDEXGROGLSIFSLAARGICIAHE 240
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DB 298 EAMASSSLIAMPREYFHVVGTTIGFLKAGQIPGFRFELQKVPKRSVHNGFAKEFWETP 357
QY 355 -----GEREOGL-----BEDVVGORPCQDCITLON 380
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QY 381 VSAGLNHQTSSVVAAYVSVAQALNHTLOQ-----NASCCPAQDPVKPQWLENNYN 432
DB 417 -----ISYVYLAAYSIHAADIDYTCIRGRGLFTNGS-CADIKKVEAWQVUKLRLH 467
QY 433 LTFHVG-GLPLRFDSGGVNDMEYDKLW---VMQSSVRLHVDYGRFN-GSLRTERL----- 483
DB 468 LNFTSNMGEQVTFDECGDLAGNYSIINMHLSPEDSIV-FKEVGYVYAKKGERLFTND 526
QY 484 -KIRWHTSDNOKPVRCSCROCEGOVRR-VKGFHSCCYDCVDCBAGSYRQNPDDIACFTFC 541
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QY 658 IFV--ESELPLS-----WADRLSGCLRGPMWLVVLAAMLVEALCTWYLVAPPEVTD 710
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RESULT 13

US-08-353-784-5
Sequence 5, Application US/08353784
Patent No. 6011068

GENERAL INFORMATION:

APPLICANT: Edward F. Nemeth, Edward M.
APPLICANT: Brown, Steven C. Hebert,
APPLICANT: Bradford C. Van Magenen, Manuel
APPLICANT: F. Balandrin, Forrest H. Fuller,
APPLICANT: Eric G. Delmar, and Scott T. Moe
TITLE OF INVENTION: CALCIUM RECEPTOR-ACTIVE
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:

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STREET: Suite 4700
STREET: 633 West Fifth Street
CITY: Los Angeles
STATE: California
COUNTRY: USA
ZIP: 90071
COMPUTER READABLE FORM:

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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: FASTSEQ
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/353, 784
FILING DATE: 9 December, 1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA: including application
PRIOR APPLICATION DATA: described below: 8
APPLICATION NUMBER: PCT/US/94/12117
FILING DATE: 21 October, 1994
APPLICATION NUMBER: U.S. 08/292,827
FILING DATE: 23 August, 1994
APPLICATION NUMBER: U.S. 08/141,248
FILING DATE: 22 October, 1993
APPLICATION NUMBER: U.S. 08/009,389
FILING DATE: 23 February, 1993
APPLICATION NUMBER: U.S. 08/017,127
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APPLICATION NUMBER: U.S. 07/934,161
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APPLICATION NUMBER: U.S. 07/834,044
FILING DATE: 11 February, 1992
APPLICATION NUMBER: U.S. 07/749,451
FILING DATE: 23 August, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Hebert, Sheldon O.
REGISTRATION NUMBER: 38,179
REFERENCE/DOCKET NUMBER: 209/069
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1085 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-353-784-5

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Query Match 23.4%; Score 1060; DB 3; Length 1085;
Best Local Similarity 30.0%; Pred. No. 1.8e-97;
Matches 269; Conservative 155; Mismatches 362; Indels 112; Gaps 26;

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DB 62 IRYNFRFRWLQAIIFAEIEINSSPALLPMTLGYRIFTCNVSKALETLSTFAONKI 121
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QY 301 EAWLTSIDLVMGLPGMAOMGTVLGFLORGAQLHEFPQYV-KTHLALATDPAFCAL----- 354
DB 298 EAMASSSLIAMPREYFHVVGTTIGFLKAGQIPGFRFELQKVPKRSVHNGFAKEFWETP 357
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Search completed: May 11, 2004, 15:30:34
Job time : 22.1574 secs

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